

<110> ASAHI KASEI KABUSHIKI KAISHA
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 Goichi HONDA
 Shuji MURAMATSU
 Yukiko NAGANO

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Leu Leu Ala Gln Leu Ser Asp Ala Ala Lys Asn Phe Glu Asp Val Arg
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Cys Lys Cys Ile Cys Pro Pro Tyr Lys Glu Asn Ser Gly His Ile Tyr
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Asn Lys Asn Ile Ser Gln Lys Asp Cys Asp Cys Leu His Val Val Glu
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Pro Met Pro Val Arg Gly Pro Asp Val Glu Ala Tyr Cys Leu Arg Cys
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Pro Ser Tyr Asn Val Ala Thr Thr Leu Pro Ser Tyr Asp Glu Ala Glu 65 70 75 80

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Gly 145	Arg	Tyr	Gly	Ala	Ile 150	Ser	Gly	Phe	Gly	Leu 155	Ser	Leu	Ile	Lys	Trp 160	
Ile	Leu	Ile	Val	Arg 165	Phe	Ser	Thr	Tyr	Phe 170	Pro	Gly	Tyr	Phe	Asp 175	Gly	
Gln	Tyr	Trp	Leu 180	Trp	Trp	Val	Phe	Leu 185	Val	Leu	Gly	Phe	Leu 190	Leu	Phe	
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Asn Pro Ala Pro Gln Ile Val Gln Ala Ala Ser Ser Ala Pro Ala Leu 35 40 45

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Pro Thr Thr Ser Asp Thr Glu Val Tyr Gly Glu Phe Tyr Pro Val Pro 65 70 75 80

Pro Pro Tyr Ser Val Ala Thr Ser Leu Pro Thr Tyr Asp Glu Ala Glu 85 90 95

Lys Ala Lys Ala Ala Ala Met Ala Ala Ala Ala Ala Glu Thr Ser Gln 100 105 110

Arg Ile Gln Glu Glu Cys Pro Pro Arg Asp Asp Phe Ser Asp Ala

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Met Ala Phe Ile Phe Asn Trp Leu Gly Phe Cys Leu Ser Phe Cys Ile

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<212> PRT

<213> Homo sapiens

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- Ala Ala Thr Thr Ser Ser Thr Gly Val Ala Val Gly Ala Glu His Gly 65 70 75 80
- Glu Asp Ser Leu Ser Arg Lys Pro Asp Pro Glu Pro Gly Arg Met Asp 85 90 95
- His His Gln Pro Gly Thr Gly Arg Tyr Gln Val Leu Leu Asn Glu Glu
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- Ala Pro Gln Ile Val Gln Ala Val Ser Ser Ala Pro Ala Leu Glu Thr 130 135 140
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- Thr Ser Asp Thr Glu Val Tyr Gly Glu Phe Tyr Pro Val Pro Pro Pro 165 170 175
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- Gln Glu Glu Cys Pro Pro Arg Asp Asp Phe Ser Asp Ala Asp Gln 210 215 220
- Leu Arg Val Gly Asn Asp Gly Ile Phe Met Leu Ala Phe Phe Met Ala 225 230 235 240
- Phe Ile Phe Asn Trp Leu Gly Phe Cys Leu Ser Phe Cys Ile Thr Asn 245 250 255
- Thr Ile Ala Gly Arg Tyr Gly Ala Ile Cys Gly Phe Gly Leu Ser Leu 260 265 270
- Ile Lys Trp Ile Leu Ile Val Arg Phe Ser Asp Tyr Phe Thr Gly Tyr 275 280 285
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Ser Ala Arg Gly Ala Pro Glu Leu Leu Arg Gly Thr Ala Thr Asn Ala
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Ser Leu Ser Arg Lys Pro Asp Pro Glu Pro Gly Arg Met Asp His His
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Gln Pro Gly Thr Gly Arg Tyr Gln Val Leu Leu Asn Glu Glu Asp Asn
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Asp Ser Ser Pro Pro Pro Tyr Ser Ser Ile Thr Val Glu Val Pro Thr 145 150 155 160

Thr Ser Asp Thr Glu Val Tyr Gly Glu Phe Tyr Pro Val Pro Pro Pro 165 170 175

Tyr Ser Val Ala Thr Ser Leu Pro Thr Tyr Asp Glu Ala Glu Lys Ala 180 185 190

Lys Ala Ala Ala Met Ala Ala Ala Ala Glu Thr Ser Gln Arg Ile 195 200 205

Gln Glu Glu Cys Pro Pro Arg Asp Asp Phe Ser Asp Ala Asp Gln 210 215 220

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Phe Asn Gly Gln Tyr Trp Leu Trp Trp Ile Phe Leu Val Leu Gly Leu 290 295 300

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Lys Phe Leu Glu Pro Tyr Ile Tyr Pro Leu Val Ser Pro Phe Val Ser
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Pro Leu Phe Gly Ile Met Ser Ser Asp Ser Ala Asp Pro Val His Ala
Val Val Tyr Ile Val Phe Met Leu Gly Ser Cys Ala Phe Phe Ser Lys
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Val His Glu Leu Asn Arg Tyr Ile Pro Thr Ala Ala Ala Phe Gly Gly
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                                                           Met
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Tyr Asn Phe Ser Leu Val Ala Leu Ser Leu Tyr Ile Val Tyr Glu Phe 65 70 75 80

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Asp Tyr Ser Asn Ser Pro Glu Ala Leu Arg Met Val Arg Val Ala Trp
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Leu Phe Leu Phe Ser Lys Phe Ile Glu Leu Met Asp Thr Val Ile Phe 115 120 125

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Tyr Lys Thr Cys Arg Arg Pro Arg Pro Val Val Thr Thr Thr Ser

Thr Thr Val Val His Ala Pro Tyr Pro Gln Pro Pro Ser Val Pro Pro 50 60

Ser Tyr Pro Gly Pro Ser Tyr Gln Gly Tyr His Thr Met Pro Pro Gln 65 70 75 80

Pro Gly Met Pro Ala Ala Pro Tyr Pro Met Gln Tyr Pro Pro Pro Tyr 85 90 95

Pro Ala Gln Pro Met Gly Pro Pro Ala Tyr His Glu Thr Leu Ala Gly
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Gly Ala Ala Ala Pro Tyr Pro Ala Ser Gln Pro Pro Tyr Asn Pro Ala 115 120 125

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Tyr Lys Thr Cys Arg Arg Pro Arg Pro Val Val Thr Thr Thr Ser 35 40 45

Thr Thr Val Val His Ala Pro Tyr Pro Gln Pro Pro Ser Val Pro Pro 50 55 60

Ser Tyr Pro Gly Pro Ser Tyr Gln Gly Tyr His Thr Met Pro Pro Gln 65 70 75 80

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85

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<212> PRT

<213> Homo sapiens

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Phe 65	Pro	Gly	Ala	Ser	Leu 70	Tyr	Leu	Pro	Met	Ala 75	Gln	Ser	Val	Ala	Val 80	
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<211> 455

<212> PRT

<213> Homo sapiens

<400> 35

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Tyr Glu Ile Arg Gln Tyr Val Val Gln Val Ile Phe Ser Val Thr Phe 35 40 45

Ala Phe Ser Cys Thr Met Phe Glu Leu Ile Ile Phe Glu Ile Leu Gly 50 55 60

Val Leu Asn Ser Ser Ser Arg Tyr Phe His Trp Lys Met Asn Leu Cys
65 70 75 80

Val Ile Leu Leu Ile Leu Val Phe Met Val Pro Phe Tyr Ile Gly Tyr 85 90 95

Phe Ile Val Ser Asn Ile Arg Leu Leu His Lys Gln Arg Leu Leu Phe 100 105 110

Ser Cys Leu Leu Trp Leu Thr Phe Met Tyr Phe Phe Trp Lys Leu Gly 115 120 125

Asp Pro Phe Pro Ile Leu Ser Pro Lys His Gly Ile Leu Ser Ile Glu 130 135 140

Leu Ser Gly Phe Gly Ala Val Asn Cys Pro Tyr Thr Tyr Met Ser Tyr 165 170 175

Phe Leu Arg Asn Val Thr Asp Thr Asp Ile Leu Ala Leu Glu Arg Arg

180 185 190 Leu Leu Gln Thr Met Asp Met Ile Ile Ser Lys Lys Arg Met Ala 200 Met Ala Arg Arg Thr Met Phe Gln Lys Gly Glu Val His Asn Lys Pro 215 Ser Gly Phe Trp Gly Met Ile Lys Ser Val Thr Thr Ser Ala Ser Gly Ser Glu Asn Leu Thr Leu Ile Gln Gln Glu Val Asp Ala Leu Glu Glu Leu Ser Arg Gln Leu Phe Leu Glu Thr Ala Asp Leu Tyr Ala Thr Lys Glu Arg Ile Glu Tyr Ser Lys Thr Phe Lys Gly Lys Tyr Phe Asn Phe 280 Leu Gly Tyr Phe Phe Ser Ile Tyr Cys Val Trp Lys Ile Phe Met Ala Thr Ile Asn Ile Val Phe Asp Arg Val Gly Lys Thr Asp Pro Val Thr Arg Gly Ile Glu Ile Thr Val Asn Tyr Leu Gly Ile Gln Phe Asp Val 325 330 Lys Phe Trp Ser Gln His Ile Ser Phe Ile Leu Val Gly Ile Ile Ile 345 Val Thr Ser Ile Arg Gly Leu Leu Ile Thr Leu Thr Lys Phe Phe Tyr Ala Ile Ser Ser Ser Lys Ser Ser Asn Val Ile Val Leu Leu Leu Ala Gln Ile Met Gly Met Tyr Phe Val Ser Ser Val Leu Leu Ile Arg Met Ser Met Pro Leu Glu Tyr Arg Thr Ile Ile Thr Glu Val Leu Gly Glu Leu Gln Phe Asn Phe Tyr His Arg Trp Phe Asp Val Ile Phe Leu Val Ser Ala Leu Ser Ser Ile Leu Phe Leu Tyr Leu Ala His Lys Gln Ala Pro Glu Lys Gln Met Ala Pro 450

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														gaa Glu		550
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tct Ser	gga Gly	ttt Phe	ggt Gly 165	gct Ala	gtc Val	aac Asn	tgc Cys	cca Pro 170	tac Tyr	act Thr	tac Tyr	atg Met	tct Ser 175	tac Tyr	ttc Phe	646

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										aag Lys						742
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										act Thr						838
										gat Asp						886
										cta Leu						934
										aaa Lys						982
										aaa Lys 300						1030
										acg Thr						1078
										atc Ile						1126
										gtt Val						1174
aca Thr	tcc Ser 355	Ile	aga Arg	gga Gly	ttg Leu	ctg Leu 360	Ile	act Thr	ctt Leu	acc Thr	aag Lys 365	Phe	ttt Phe	tat Tyr	gcc Ala	1222
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					Phe					ctg Leu					Ser	1318
atg	cct	tta	gaa	tac	cgc	acc	ata	atc	act	gaa	gto	ctt	gga	gaa	ctg	1366

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Leu Ser Lys Ala Glu Arg Ala Thr Glu Leu Ala Ala Glu Gly Gln Leu 145 150 155 160

Thr Leu Gln Gln Phe Ala Gln Ser Thr Glu Met Leu Lys Arg Val Val 165 170 175

Gln Glu His Leu Pro Leu Met Ser Glu Ala Gly Ala Gly Leu Pro Asp \$180\$

Met Glu Ala Val Ala Gly Ala Glu Ala Leu Asn Gly Gln Ser Asp Phe 195 200 205

Pro Tyr Leu Gly Ala Phe Pro Ile Asn Pro Gly Leu Phe Ile Met Thr 210 215 220

Pro Ala Gly Val Phe Leu Ala Glu Ser Ala Leu His Met Ala Gly Leu 225 230 235 240

Ala Glu Tyr Pro Met Gln Gly Glu Leu Ala Ser Ala Ile Ser Ser Gly 245 250 255

Lys Lys Lys Arg Lys Arg Cys Gly Met Cys Ala Pro Cys Arg Arg Arg 260 265 270

Ile Asn Cys Glu Gln Cys Ser Ser Cys Arg Asn Arg Lys Thr Gly His

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			agt Ser													441
			cca Pro													489
			aac Asn 70													537
			ggc Gly													585
			gct Ala													633
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			tca Ser		His											729
			cgg Arg 150													777
			gcg Ala													825
			ctg Leu													873
~ ~ +	~+~	~~~	aat	~~~	~~~	~~~	ata	2.2.±	aaa	g 2 g	tee	a20	++0	ccc	tac	921

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ggt gt Gly Va	tg ttc al Phe	ctg Leu 230	gcc Ala	gag Glu	agc Ser	gcg Ala	ctg Leu 235	cac His	atg Met	gcg Ala	ggc Gly	ctg Leu 240	gct Ala	gag Glu	1017
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Ile Glu Asp Asn Glu Leu Asn Ile Val Leu Glu Leu Ala Asp Ala Gly 115 120

Asp Leu Ser Gln Met Ile Lys Tyr Phe Lys Lys Gln Lys Arg Leu Ile 130 135 140

Glu His Met His Ser Arg Arg Val Met His Arg Asp Ile Lys Pro Ala 165 170 175

Asn Val Phe Ile Thr Ala Thr Gly Val Val Lys Leu Gly Asp Leu Gly 180 185 190

Leu Gly Arg Phe Phe Ser Ser Glu Thr Thr Ala Ala His Ser Leu Val 195 200 205

Gly Thr Pro Tyr Tyr Met Ser Pro Glu Arg Ile His Glu Asn Gly Tyr 210 220

Asn Phe Lys Ser Asp Ile Trp Ser Leu Gly Cys Leu Leu Tyr Glu Met 225 230 235 240

Ala Ala Leu Gln Ser Pro Phe Tyr Gly Asp Lys Met Asn Leu Phe Ser 245 250 255

Leu Cys Gln Lys Ile Glu Gln Cys Asp Tyr Pro Pro Leu Pro Gly Glu 260 265 270

His Tyr Ser Glu Lys Leu Arg Glu Leu Val Ser Met Cys Ile Cys Pro 275 280 285

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gtg cat Val His 25												269
tgc tcg Cys Ser 40												317
ttc agc Phe Ser												365
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		aag gag Lys Glu										461
		tat ttg Tyr Leu										509
		ttg gct Leu Ala 125										557
		cag aag Gln Lys 140			-				-		_	605
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		ctc ggt Leu Gly										749

	18	85					190					195					
_	.u Th			_	-				-						atg Met		797.
		_										-		-	atc Ile 230		845
															ccc Pro		893
		ĺу													gag Glu		941
	s A													_	tta Leu	_	989
Ğ.		_	_	_	_	_		_		-			_	-	cct Pro	_	1037
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<211> 371

<212> PRT

<213> Homo sapiens

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Pro 65	Gln	Gly	Gly	Tyr	Pro 70	Gln	Gly	Pro	Tyr	Pro 75	Gln	Gly	Gly	Tyr	Pro 80
Gln	Gly	Pro	Tyr	Pro 85	Gln	Glu	Gly	Tyr	Pro 90	Gln	Gly	Pro	Tyr	Pro 95	Gln
Gly	Gly	Tyr	Pro 100	Gln	Gly	Pro	Tyr	Pro 105	Gln	Ser	Pro	Phe	Pro 110	Pro	Asn
Pro	Tyr	Gly 115	Gln	Pro	Gln	Val	Phe 120	Pro	Gly	Gln	Asp	Pro 125	Asp	Ser	Pro
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Gln 145	Asp	Phe	Pro	Ala	Thr 150	Asn	Trp	Asp	Asp	Lys 155	Ser	Ile	Arg	Gln	Ala 160
Phe	Ile	Arg	Lys	Val 165	Phe	Leu	Val	Leu	Thr 170	Leu	Gln	Leu	Ser	Val 175	Thr
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Ala	Val	Gly	Ile 260	Thr	Thr	Ala	Val	Cys 265	Phe	Thr	Val	Val	Ile 270	Phe	Ser
Met	Gln	Thr 275	Arg	Tyr	Asp	Phe	Thr 280	Ser	Cys	Met	Gly	Val 285	Leu	Leu	Val
Ser	Met 290	Val	Val	Leu	Phe	Ile 295	Phe	Ala	Ile	Leu	Cys 300	Ile	Phe	Ile	Arg
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	_			_				- ,	_					ggg	-	210
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	ccc Pro															546
	gac Asp															594
	acc Thr 170															642
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	gtc Val															738
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	ttc Phe					Phe										930
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	att Ile													Val		1026
	tca Ser															1074
	ctg Leu 330											Pro				1122

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Arg Cys Leu Val Leu Thr Gly Phe Gly Gly Tyr Asp Lys Val Lys Leu

 ${\tt Gln \ Ser \ Arg \ Pro \ Ala \ Pro \ Pro \ Ala \ Pro \ Gly \ Pro \ Gly \ Gln \ Leu \ Thr}$

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Lys Ala Gly Asp Arg Val Met Val Leu Asn Arg Ser Gly Met Trp Gln 130 135 140

Glu Glu Val Thr Val Pro Ser Val Gln Thr Phe Leu Ile Pro Glu Ala 145 150 155 160

Met Thr Phe Glu Glu Ala Ala Ala Leu Leu Val Asn Tyr Ile Thr Ala 165 170 175

Tyr Met Val Leu Phe Asp Phe Gly Asn Leu Gln Pro Gly His Ser Val 180 185 190

Leu Val His Met Ala Ala Gly Gly Val Gly Met Ala Ala Val Gln Leu 195 200 205

Cys Arg Thr Val Glu Asn Val Thr Val Phe Gly Thr Ala Ser Ala Ser 210 215 220

Lys His Glu Ala Leu Lys Glu Asn Gly Val Thr His Pro Ile Asp Tyr 225 230 235 235 240

His Thr Thr Asp Tyr Val Asp Glu Ile Lys Lys Ile Ser Pro Lys Gly
245 250 255

Val Asp Ile Val Met Asp Pro Leu Gly Gly Ser Asp Thr Ala Lys Gly 260 265 270

Tyr Asn Leu Leu Lys Pro Met Gly Lys Val Val Thr Tyr Gly Met Ala 275 280 285

Asn Leu Leu Thr Gly Pro Lys Arg Asn Leu Met Ala Leu Ala Arg Thr 290 295 300

Trp Trp Asn Gln Phe Ser Val Thr Ala Leu Gln Leu Leu Gln Ala Asn 305 310 315 320

Arg Ala Val Cys Gly Phe His Leu Gly Tyr Leu Asp Gly Glu Val Glu 325 330 335

Leu Val Ser Gly Val Val Ala Arg Leu Leu Ala Leu Tyr Asn Gl
n Gly 340 345 350

His Ile Lys Pro His Ile Asp Ser Val Trp Pro Phe Glu Lys Val Ala 355 360 365

Asp Ala Met Lys Gln Met Gln Glu Lys Lys Asn Val Gly Lys Val Leu 370 375 380

Leu Val Pro Gly Pro Glu Lys Glu Asn 385 - 390

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Arg Cys Leu Val Leu Thr Gly Phe Gly Gly Tyr Asp Lys Val Lys Leu 50 60

Gln Ser Arg Pro Ala Ala Pro Pro Ala Pro Gly Pro Gly Gln Leu Thr .65 70 75 80

Leu Arg Leu Arg Ala Cys Gly Leu Asn Phe Ala Asp Leu Met Ala Arg 90 Gln Gly Leu Tyr Asp Arg Leu Pro Pro Leu Pro Val Thr Pro Gly Met 105 Glu Gly Ala Gly Val Val Ile Ala Val Gly Glu Gly Val Ser Asp Arg 120 Lys Ala Gly Asp Arg Val Met Val Leu Asn Arg Ser Gly Met Trp Gln Glu Glu Val Thr Val Pro Ser Val Gln Thr Phe Leu Ile Pro Glu Ala 155 Met Thr Phe Glu Glu Ala Ala Ala Leu Leu Val Asn Tyr Ile Thr Ala 170 Tyr Met Val Leu Phe Asp Phe Gly Asn Leu Gln Pro Gly His Ser Val Leu Val His Met Ala Ala Gly Gly Val Gly Met Ala Ala Val Gln Leu Cys Arg Thr Val Glu Asn Val Thr Val Phe Gly Thr Ala Ser Ala Ser 210 Lys His Glu Ala Leu Lys Glu Asn Gly Val Thr His Pro Ile Asp Tyr His Thr Thr Asp Tyr Val Asp Glu Ile Lys Lys Ile Ser Pro Lys Gly 245 Val Asp Ile Val Met Asp Pro Leu Gly Gly Ser Asp Thr Ala Lys Gly Tyr Asn Leu Lys Pro Met Gly Lys Val Val Thr Tyr Gly Met Ala Asn Leu Leu Thr Gly Pro Lys Arg Asn Leu Met Ala Leu Ala Arg Thr Trp Trp Asn Gln Phe Ser Val Thr Ala Leu Gln Leu Leu Gln Ala Asn Arg Ala Val Cys Gly Phe His Leu Gly Tyr Leu Asp Gly Glu Val Glu Leu Val Ser Gly Val Val Ala Arg Leu Leu Ala Leu Tyr Asn Gln Gly His Ile Lys Pro His Ile Asp Ser Val Trp Pro Phe Glu Lys Val Ala 360 Asp Ala Met Lys Gln Met Gln Glu Lys Lys Asn Val Gly Lys Val Leu 370 375 380

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L			_										gtc Val				634
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<213> Homo sapiens

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Pro	Val	Leu 115	Gly	Ser	Leu	Leu	Asn 120	Leu	Pro	Gly	Ile	Arg 125	Ser	Phe	Val	
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	ttc Phe															214
	att Ile															262
	gaa Glu															310
	ggt Gly							Phe					Gly			358

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Ala Leu Leu Asn Asp Leu Lys Lys His Thr Ala Asp Glu Asn Pro Asp Lys Ser Thr Leu Glu Lys Ala Ile Gly Ser Leu Lys Glu Val Met Thr His Ile Asn Glu Asp Lys Arg Lys Thr Glu Ala Gln Lys Gln Ile Phe Asp Val Val Tyr Glu Val Asp Gly Cys Pro Ala Asn Leu Leu Ser 105 Ser His Arg Ser Leu Val Gln Arg Val Glu Thr Ile Ser Leu Gly Glu 120 His Pro Cys Asp Arg Gly Glu Gln Val Thr Leu Phe Leu Phe Asn Asp 135 Cys Leu Glu Ile Ala Arg Lys Arg His Lys Val Ile Gly Thr Phe Arg Ser Pro His Gly Gln Thr Arg Pro Pro Ala Ser Leu Lys His Ile His Leu Met Pro Leu Ser Gln Ile Lys Lys Val Leu Asp Ile Arg Glu Thr Glu Asp Cys His Asn Ala Phe Ala Leu Leu Val Arg Pro Pro Thr Glu 200 Gln Ala Asn Val Leu Leu Ser Phe Gln Met Thr Ser Asp Glu Leu Pro Lys Glu Asn Trp Leu Lys Met Leu Cys Arg His Val Ala Asn Thr Ile 230 235 Cys Lys Ala Asp Ala Glu Asn Leu Ile Tyr Thr Ala Asp Pro Glu Ser Phe Glu Val Asn Thr Lys Asp Met Asp Ser Thr Leu Ser Arg Ala Ser Arg Ala Ile Lys Lys Thr Ser Lys Lys Val Thr Arg Ala Phe Ser Phe Ser Lys Thr Pro Lys Arg Ala Leu Arg Arg Ala Leu Met Thr Ser His Gly Ser Val Glu Gly Arg Ser Pro Ser Ser Asn Asp Lys His Val Met 315 Ser Arg Leu Ser Ser Thr Ser Ser Leu Ala Gly Ile Pro Ser Pro Ser 330

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Glu	Thr	Ile	Ser 125	Leu	Gly	Glu	His	Pro 130	Cys	Asp	Arg	Gly	Glu 135	Gln	Val	
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_	gtt Val 155						_					Thr	_			833
_	tct Ser															881
	ttg Leu															929
	gtg Val						_	_					_		-	977
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_	cat His 235	_	_				_		-	-	-					1073
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	gct Ala															1265
_	aat Asn 315	-	_		-	_	-	_			_					1313
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350 355

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<213> Homo sapiens

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Gln Ser Lys His Met Lys Gln Thr Leu Ser Leu Arg Lys Thr Pro Tyr 565 570 575

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		tgg Trp														1699
		aca Thr 535														1747
ggc Gly	ctg Leu 550	acc Thr	tcc Ser	cat His	gag Glu	aga Arg 555	atc Ile	agc Ser	ctg Leu	cag Gln	aag Lys 560	cag Gln	agc Ser	aag Lys	cat His	1795
		cag Gln														1843
atg Met	cag Gln	aac Asn	ctg Leu	gca Ala 585	gat Asp	ttc Phe	ttt Phe	cag Gln	tgt Cys 590	ggc Gly	tgc Cys	ttt Phe	ggc Gly	ttg Leu 595	gtg Val	1891
aag Lys	ccc Pro	tgt Cys	gtg Val 600	gta Val	gat Asp	tgg Trp	aca Thr	tca Ser 605	cag Gln	tac Tyr	acc Thr	atg Met	gtc Val 610	ttt Phe	cac His	1939
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Asp Ile Pro Val Pro Tyr Leu Tyr Phe Asp Met Gly Ala Ala Val Leu 50 60

Cys Ala Ser Phe Met Ser Phe Gly Val Lys Arg Arg Trp Phe Ala Leu 65 70 75 80

Gly Ala Ala Leu Gln Leu Ala Ile Ser Thr Tyr Ala Ala Tyr Ile Gly 85 90 95

Gly Tyr Val His Tyr Gly Asp Trp Leu Lys Val Arg Met Tyr Ser Arg 100 105 110

Thr Val Ala Ile Ile Gly Gly Phe Leu Val Leu Ala Ser Gly Ala Gly 115 120 125

Glu Leu Tyr Arg Arg Lys Pro Arg Ser Arg Ser Leu Gln Ser Thr Gly 130 135

Gln Val Phe Leu Gly Ile Tyr Leu Ile Cys Val Ala Tyr Ser Leu Gln 145 150 155 160

His Ser Lys Glu Asp Arg Leu Ala Tyr Leu Asn His Leu Pro Gly Gly 165 170 175

Glu Leu Met Ile Gln Leu Phe Phe Val Leu Tyr Gly Ile Leu Ala Leu 180 185 190

Ala Phe Leu Ser Gly Tyr Tyr Val Thr Leu Ala Ala Gln Ile Leu Ala 195 200 205

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Val Leu Leu Pro Pro Val Met Leu Leu Ile Asp Gly Asn Val Ala Tyr
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Trp His Asn Thr Arg Arg Val Glu Phe Trp Asn Gln Met Lys Leu Leu
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Ile Met Gln Leu Gly Ser Val Leu Leu Thr Arg Cys Pro Phe Trp Gly
tgc ttc agc cag ctc atg ctg tac gct gag agg gct gag gca cgc cgg
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Cys Phe Ser Gln Leu Met Leu Tyr Ala Glu Arg Ala Glu Ala Arg Arg
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aag ccc gac atc cca gtg cct tac ctg tat ttc gac atg ggg gca gcc
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Lys Pro Asp Ile Pro Val Pro Tyr Leu Tyr Phe Asp Met Gly Ala Ala
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Val Leu Cys Ala Ser Phe Met Ser Phe Gly Val Lys Arg Arg Trp Phe
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gcg ctg ggg gcc gca ctc caa ttg gcc att agc acc tac gcc gcc tac
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atc ggg ggc tac gtc cac tac ggg gac tgg ctg aag gtc cgt atg tac
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teg ege aca gtt gee ate ate gge gge ttt ett gtg ttg gee age ggt
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Ser Arg Thr Val Ala Ile Ile Gly Gly Phe Leu Val Leu Ala Ser Gly
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Ala Gly Glu Leu Tyr Arg Arg Lys Pro Arg Ser Arg Ser Leu Gln Ser
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Gln Ile Gln Lys Glu Gln Asp Tyr Gln Arg Tyr Arg Glu Glu Arg Phe
Arg Leu Thr Ser Glu Ser Thr Asn Gln Arg Val Leu Trp Trp Ser Ile
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Ile Gln Val Gly Glu His Ala Asn Asn Tyr Pro Glu Ile Ala Ala Lys
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Asp Lys Leu Thr Glu Leu Gln Leu Arg Ala Arg Gln Leu Leu Asp Gln
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cgt cac ct Arg His Le 95	u Lys Ser P	tc ttt gag he Phe Glu 00	gcc aag aag Ala Lys Lys 105	ctg gtg tag Leu Val	gtgccctc	339
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<213> Homo sapiens

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cat His	gtt Val	cca Pro	cct Pro	tcc Ser 230	ctc Leu	ttc Phe	acg Thr	gaa Glu	cct Pro 235	aat Asn	gaa Glu	ata Ile	tcc Ser	cag Gln 240	tat Tyr	839
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Lys Lys Ile Ser Arg Leu Asp Ala Glu Leu Val Lys Tyr Lys Asp Gln 35 40 45

Ile Lys Lys Met Arg Glu Gly Pro Ala Lys Asn Met Val Lys Gln Lys 50 55 60

Ala Leu Arg Val Leu Lys Gln Lys Arg Met Tyr Glu Gln Gln Arg Asp
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Asn Leu Ala Gln Gln Ser Phe Asn Met Glu Gln Ala Asn Tyr Thr Ile $85 \hspace{1cm} 90 \hspace{1cm} 95$

Gln Ser Leu Lys Asp Thr Lys Thr Thr Val Asp Ala Met Lys Leu Gly
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Val Lys Glu Met Lys Lys Ala Tyr Lys Gln Val Lys Ile Asp Gln Ile 115 120 125

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Ile Gln Glu Ala Leu Ser Arg Ser Tyr Gly Thr Pro Glu Leu Asp Glu 145 150 155 160

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					gca Ala											249
					agg Arg											297
					atg Met											345
					acg Thr 105											393
					aag Lys											441
					gat Asp											489
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Trp Asp Ile Val Lys Ala Thr Gln Tyr Gly Ile Tyr Glu Arg Cys Arg 50 55 60

Glu Leu Val Glu Ala Gly Tyr Asp Val Arg Gln Pro Asp Lys Glu Asn
65 70 75 80

Val Thr Leu Leu His Trp Ala Ala Ile Asn Asn Arg Ile Asp Leu Val 85 90 95

Lys Tyr Tyr Ile Ser Lys Gly Ala Ile Val Asp Gln Leu Gly Gly Asp
100 105 110

Leu Asn Ser Thr Pro Leu His Trp Ala Thr Arg Gln Gly His Leu Ser

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Gly 145	Glu	Gly	Cys	Ser	Cys 150	Ile	His	Leu	Ala	Ala 155	Gln	Phe	Gly	His	Thr 160
Ser	Ile	Val	Ala	Tyr 165	Leu	Ile	Ala	Lys	Gly 170	Gln	Asp	Val	Asp	Met 175	Met
Asp	Gln	Asn	Gly 180	Met	Thr	Pro	Leu	Met 185	Trp	Ala	Ala	Tyr	Arg 190	Thr	His
Ser	Val	Asp 195	Pro	Thr	Arg	Leu	Leu 200	Leu	Thr	Phe	Asn	Val 205	Ser	Val	Asn
Leu	Gly 210	Asp	Lys	Tyr	His	Lys 215	Asn	Thr	Ala	Leu	His 220	Trp	Ala	Val	Leu
Ala 225	Gly	Asn	Thr	Thr	Val 230	Ile	Ser	Leu	Leu	Leu 235	Glu	Ala	Gly	Ala	Asn 240
Val	Asp	Ala	Gln	Asn 245	Ile	Lys	Gly	Glu	Ser 250	Ala	Leu	Asp	Leu	Ala 255	Lys
Gln	Arg	Lys	Asn 260	Val	Trp	Met	Ile	Asn 265	His	Leu	Gln	Glu	Ala 270	Arg	Gln
Ala	Lys	Gly 275	Tyr	Asp	Asn	Pro	Ser 280	Phe	Leu	Arg	Lys	Leu 285	Lys	Ala	Asp
Lys	Glu 290	Phe	Arg	Gln	Lys	Val 295	Met	Leu	Gly	Thr	Pro 300	Phe	Leu	Val	Ile
Trp 305	Leu	Val	Gly	Phe	Ile 310	Ala	Asp	Leu	Asn	Ile 315	Asp	Ser	Trp	Leu	Ile 320
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Lys	Ser	Phe	Phe 340	Asp	His	Ser	Met	His 345	Ser	Ala	Leu	Pro	Leu 350	Gly	Ile
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Trp	Asn 370	Asp	Leu	Asn	Phe	Leu 375	Phe	Ile	His	Leu	Pro 380	Phe	Leu	Ala	Asn
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Gly	Ile	Ile	Lys	Ala 405	Thr	Glu	Glu	Gln	Lys 410	Lys	Lys	Thr	Ile	Val 415	Glu
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Cys	Ile 450	Ala	Lys	Phe	Asp	His 455	His	Cys	Pro	Trp	Val 460	Gly	Asn	Cys	Val	
Gly 465	Ala	Gly	Asn	His	Arg 470	Tyr	Phe	Met	Gly	Tyr 475	Leu	Phe	Phe	Leu	Leu 480	
Phe	Met	Ile	Cys	Trp 485	Met	Ile	Tyr	Gly	Cys 490	Ile	Ser	Tyr	Trp	Gly 495	Leu	
His	Cys	Glu	Thr 500	Thr	Tyr	Thr	Lys	Asp 505	Gly	Phe	Trp	Thr	Tyr 510	Ile	Thr	
Gln	Ile	Ala 515	Thr	Cys	Ser	Pro	Trp 520	Met	Phe	Trp	Met	Phe 525	Leu	Asn	Ser	
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Cys	Gly	Leu 595		Arg	Pro	Val	Ile 600		Asp	Trp	Thr	Arg 605	Gln	Туг	Thr	
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ggc Gly	tgt Cys	z gto s Va.	l Pro	c cti	t cto u Leo	c cad	c cca s Pro	o Gl	g gaa ı Glı	a ato u Ilo	c aaa e Lya	a cco s Pro 2.	o Gli	a age n Se:	c cat r His	97

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gat Asp 45	tac Tyr	agc Ser	aca Thr	tgg Trp	gac Asp 50	ata Ile	gtc Val	aag Lys	gct Ala	aca Thr 55	caa Gln	tat Tyr	gga Gly	ata Ile	tat Tyr 60	193
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Arg Lys Thr His Ile Asp Asp Tyr Ser Thr Trp Asp Ile Val Lys Ala 50 55 60

Thr Gln Tyr Gly Ile Tyr Glu Arg Cys Arg Glu Leu Val Glu Ala Gly
65 70 75 80

Tyr Asp Val Arg Gln Pro Asp Lys Glu Asn Val Thr Leu Leu His Trp 85 90 95

Ala Ala Ile Asn Asn Arg Ile Asp Leu Val Lys Tyr Tyr Ile Ser Lys 100 105 110

Gly Ala Ile Val Asp Gln Leu Gly Gly Asp Leu Asn Ser Thr Pro Leu 115 120 125

His Trp Ala Thr Arg Gln Gly His Leu Ser Met Val Val Gln Leu Met 130 135 140

Lys Tyr Gly Ala Asp Pro Ser Leu Ile Asp Gly Glu Gly Cys Ser Cys 145 150 155 160

Ile His Leu Ala Ala Gln Phe Gly His Thr Ser Ile Val Ala Tyr Leu 165 170 175

Ile Ala Lys Gly Gln Asp Val Asp Met Met Asp Gln Asn Gly Met Thr 180 185 190 Pro Leu Met Trp Ala Ala Tyr Arg Thr His Ser Val Asp Pro Thr Arg Leu Leu Leu Thr Phe Asn Val Ser Val Asn Leu Gly Asp Lys Tyr His 215 Lys Asn Thr Ala Leu His Trp Ala Val Leu Ala Gly Asn Thr Thr Val Ile Ser Leu Leu Glu Ala Gly Ala Asn Val Asp Ala Gln Asn Ile Lys Gly Glu Ser Ala Leu Asp Leu Ala Lys Gln Arg Lys Asn Val Trp 265 Met Ile Asn His Leu Gln Glu Ala Arg Gln Ala Lys Gly Tyr Asp Asn Pro Ser Phe Leu Arg Lys Leu Lys Ala Asp Lys Glu Phe Arg Gln Lys Val Met Leu Gly Thr Pro Phe Leu Val Ile Trp Leu Val Gly Phe Ile 310 315 Ala Asp Leu Asn Ile Asp Ser Trp Leu Ile Lys Gly Leu Met Tyr Gly 325 330 Gly Val Trp Ala Thr Val Gln Phe Leu Ser Lys Ser Phe Phe Asp His 345 Ser Met His Ser Ala Leu Pro Leu Gly Ile Tyr Leu Ala Thr Lys Phe Trp Met Tyr Val Thr Trp Phe Phe Trp Phe Trp Asn Asp Leu Asn Phe Leu Phe Ile His Leu Pro Phe Leu Ala Asn Ser Val Ala Leu Phe Tyr Asn Phe Gly Lys Ser Trp Lys Ser Asp Pro.Gly Ile Ile Lys Ala. Thr 410 Glu Glu Gln Lys Lys Thr Ile Val Glu Leu Ala Glu Thr Gly Ser Leu Asp Leu Ser Ile Phe Cys Ser Thr Cys Leu Ile Arg Lys Pro Val Arg Ser Lys His Cys Gly Val Cys Asn Arg Cys Ile Ala Lys Phe Asp 455 His His Cys Pro Trp Val Gly Asn Cys Val Gly Ala Gly Asn His Arg Tyr Phe Met Gly Tyr Leu Phe Phe Leu Leu Phe Met Ile Cys Trp Met

485

490

Ile Tyr Gly Cys Ile Ser Tyr Trp Gly Leu His Cys Glu Thr Thr Tyr 500 505 Thr Lys Asp Gly Phe Trp Thr Tyr Ile Thr Gln Ile Ala Thr Cys Ser 520 Pro Trp Met Phe Trp Met Phe Leu Asn Ser Val Phe His Phe Met Trp Val Ala Val Leu Leu Met Cys Gln Met Tyr Gln Ile Ser Cys Leu Gly 550 Ile Thr Thr Asn Glu Arg Met Asn Ala Arg Arg Tyr Lys His Phe Lys 570 Val Thr Thr Thr Ser Ile Glu Ser Pro Phe Asn His Gly Cys Val Arg 585 Asn Ile Ile Asp Phe Phe Glu Phe Arg Cys Cys Gly Leu Phe Arg Pro 600 Val Ile Val Asp Trp Thr Arg Gln Tyr Thr Ile Glu Tyr Asp Gln Ile 615 Ser Gly Ser Gly Tyr Gln Leu Val 630 625 <210> 66 <211> 4715 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (108)..(2003) <400> 66 gaagaaggag gaggaggccc gcgtcgcctc cggcggggct cgcgctcgcc ccgcgctcgc 60 cetecgeete geeegageee egggagggtg aaacgettte teecage atg eag egg Met Gln Arg 1 gag gag gga ttt aac acc aag atg gcg gac ggc ccg gat gag tac gat 164 Glu Glu Gly Phe Asn Thr Lys Met Ala Asp Gly Pro Asp Glu Tyr Asp 10 5 acc gaa gcg ggc tgt gtg ccc ctt ctc cac cca gag gaa atc aaa ccc 212 Thr Glu Ala Gly Cys Val Pro Leu Leu His Pro Glu Glu Ile Lys Pro 25 20 caa agc cat tat aac cat gga tat ggt gaa cct ctt gga cgg aaa act 260 Gln Ser His Tyr Asn His Gly Tyr Gly Glu Pro Leu Gly Arg Lys Thr 45 40 308 cat att gat gat tac agc aca tgg gac ata gtc aag gct aca caa tat

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Val Arg Pro Phe Asp Gly Asn Ser Pro Ile Leu Tyr Tyr Ile Val Glu 35 40 45

Leu Ser Glu Asn Asn Ser Pro Trp Lys Val His Leu Ser Asn Val Gly 50 55 60

Pro Glu Met Thr Gly Val Thr Val Ser Gly Leu Thr Pro Ala Arg Thr 65 70 75 80

Tyr Gln Phe Arg Val Cys Ala Val Asn Glu Val Gly Arg Gly Gln Tyr 85 90 95

Ser Ala Glu Thr Ser Arg Leu Met Leu Pro Glu Glu Pro Pro Ser Ala 100 105 110

Pro Pro Lys Asn Ile Val Ala Ser Gly Arg Thr Asn Gln Ser Ile Met 115 120 125

Val Gln Trp Gln Pro Pro Pro Glu Thr Glu His Asn Gly Val Leu Arg 130 135 140

Gly Tyr Ile Leu Arg Tyr Arg Leu Ala Gly Leu Pro Gly Glu Tyr Gln 145 150 155 160

Gln Arg Asn Ile Thr Ser Pro Glu Val Asn Tyr Cys Leu Val Thr Asp 165 170 175

Leu Ile Ile Trp Thr Gln Tyr Glu Ile Gln Val Ala Ala Tyr Asn Gly
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Ala Gly Leu Gly Val Phe Ser Arg Ala Val Thr Glu Tyr Thr Leu Gln
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Gly Val Pro Thr Ala Pro Pro Gln Asn Val Gln Thr Glu Ala Val Asn 210 215 220

Ser Thr Thr Ile Gln Phe Leu Trp Asn Pro Pro Pro Gln Gln Phe Ile 225 230 235 240

Asn Gly Ile Asn Gln Gly Tyr Lys Leu Leu Ala Trp Pro Ala Asp Ala 245 250 255

Pro Glu Ala Val Thr Val Val Thr Ile Ala Pro Asp Phe His Gly Val 260 265 270

His His Gly His Ile Thr Asn Leu Lys Lys Phe Thr Ala Tyr Phe Thr 275 . 280 285

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Gln Leu Val Trp Thr Gln Glu Asp Lys Pro Gly Ala Val Gly His Leu 305 310 315 320

Ser Phe Thr Glu Ile Leu Asp Thr Ser Leu Lys Val Ser Trp Gln Glu 325 330 335

Pro Leu Glu Lys Asn Gly Ile Ile Thr Gly Tyr Gln Ile Ser Trp Glu 340 345 350

Val Tyr Gly Arg Asn Asp Ser Arg Leu Thr His Thr Leu Asn Ser Thr 355 360 365

Met His Glu Tyr Lys Ile Gln Gly Leu Ser Ser Leu Thr Thr Tyr Thr 370 375 380

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Ser Thr Ile Ser Ser Gly Val Pro Pro Asp Leu Pro Gly Ala Pro Ser 405 410 415

Asn Leu Val Ile Ser Asn Ile Ser Pro Arg Ser Ala Thr Leu Gln Phe 420 425 430

Arg Pro Gly Tyr Asp Gly Lys Thr Ser Ile Ser Arg Trp Ile Val Glu
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Gly Gln Met Arg Pro Glu Gly Val Gly Leu Pro Ala Glu Val Thr Gln 450 455 460

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ctg act ccg gct cgt acc Leu Thr Pro Ala Arg Thr 75 80	Tyr Gln Phe Arg Val	. Cys Ala Val Asn Glu
gtg ggc agg ggc cag tac Val Gly Arg Gly Gln Tyr 95	agt gcc gag aca agc Ser Ala Glu Thr Ser 100	agg ttg atg cta cct 339 Arg Leu Met Leu Pro 105
gaa gaa cca ccc agt gct Glu Glu Pro Pro Ser Ala 110	-	
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cac aac ggg gtg ttg cgt His Asn Gly Val Leu Arg 140		
ctt ccc gga gag tac cag Leu Pro Gly Glu Tyr Glr 155 160	Gln Arg Asn Ile Thr	Ser Pro Glu Val Asn
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	ccg Pro 235	cct Pro	cag Gln	cag Gln	ttt Phe	atc Ile 240	aat Asn	ggc Gly	atc Ile	aac Asn	cag Gln 245	gga Gly	tac Tyr	aag Lys	ctt Leu	ctg Leu 250	771
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- Ser Thr Thr Ile Gln Phe Leu Trp Asn Pro Pro Pro Gln Gln Phe Ile 225 230 235 240
- Asn Gly Ile Asn Gln Gly Tyr Lys Leu Leu Ala Trp Pro Ala Asp Ala 245 250 255
- Pro Glu Ala Val Thr Val Val Thr Ile Ala Pro Asp Phe His Gly Val 260 265 270
- His His Gly His Ile Thr Asn Leu Lys Lys Phe Thr Ala Tyr Phe Thr 275 280 285
- Ser Val Leu Cys Phe Thr Thr Pro Gly Asp Gly Pro Pro Ser Thr Pro 290 295 300
- Gln Leu Val Trp Thr Gln Glu Asp Lys Pro Gly Ala Val Gly His Leu 305 310 315 320
- Ser Phe Thr Glu Ile Leu Asp Thr Ser Leu Lys Val Ser Trp Gln Glu 325 330 335
- Pro Leu Glu Lys Asn Gly Ile Ile Thr Gly Tyr Gln Ile Ser Trp Glu 340 345 350
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Pro 465	Ser	His	Glu	Ala	Gly 470	Leu	Glu	Pro	Ala	Asn 475	Leu	Gly	Ser	Leu	Trp 480	
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cto	g act	t cc r Pre	g gc	t cgt a Arc	acc Thi	c tat	caa Glr	a tto n Phe	c cgg	g gto g Val	g tgo l Cys	c gcg s Ala	g gto a Val	g aat L Asr	gaa Glu	29

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					atg Met											435
					cgt Arg											483
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					gac Asp											579
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	aac Asn 475	ctc Leu	gga Gly	agt Ser	ctg Leu	tgg Trp 480	ctg Leu	ctc Leu	agc Ser	ctg Leu	gtg Val 485	Tyr	tgg Trp	tgt Cys	tac Tyr	agc Ser 490	1491
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Phe Phe Val Gly Ala Ala Ile Leu Trp Leu Leu Cys Ser His Arg Pro 50 60

Ala Pro Gly Arg Pro Pro Thr His Asn Ala His Asn Trp Arg Leu Gly 65 70 75 80

Gln Ala Pro Ala Asn Trp Tyr Asn Asp Thr Tyr Pro Leu Ser Pro Pro 85 90 95

Gln Arg Thr Pro Ala Gly Ile Arg Tyr Arg Ile Ala Val Ile Ala Asp 100 105 110

Leu Asp Thr Glu Ser Arg Ala Gln Glu Glu Asn Thr Trp Phe Ser Tyr 115 120 125

Leu Lys Lys Gly Tyr Leu Thr Leu Ser Asp Ser Gly Asp Lys Val Ala 130 135 140

Val Glu Trp Asp Lys Asp His Gly Val Leu Glu Ser His Leu Ala Glu 145 150 155 160

Lys Gly Arg Gly Met Glu Leu Ser Asp Leu Ile Val Phe Asn Gly Lys 165 170 175

Leu Tyr Ser Val Asp Asp Arg Thr Gly Val Val Tyr Gln Ile Glu Gly
180 185 190

Ser Lys Ala Val Pro Trp Val Ile Leu Ser Asp Gly Asp Gly Thr Val 195 200 205

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atg ca Met Hi 15					_							_			216
atg ac Met Th															264
ctg ac Leu Th	_				-	-				_		_			312
ege ec Arg Pr															360
ctc gg Leu Gl															408
ccc cc Pro Pr 95				_	_					_		_	_		456
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agt ta Ser Ty								_		_	_		_	_	552
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Leu Gly Gln Trp Ile Gln Arg
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Ser	Gln	Phe	Val	Glu 85	Asn	Cys	Lys	Gly	Val 90	Ile	Gln	Arg	Leu	Thr 95	Leu
Gln	Glu	His	Lys 100	Met	Val	Trp	Asn	Arg 105	Thr	Thr	His	Leu	Trp 110	Asn	Asp
Суѕ	Ser	Lys 115	Ile	Ile	His	Gln	Arg 120	Thr	Asn	Thr	Val	Pro 125	Phe	Asp	Leu
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Phe	Ile	Leu	Arg 260	Lys	Gln	Tyr	Leu	Gln 265	Arg	Gln	Glu	Arg	Leu 270	Arg	Leu
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gag Glu	gag Glu	ttc Phe	cag Gln 280	gag Glu	cat His	gag Glu	gcc Ala	cag Gln 285	ctg Leu	ctg Leu	agc Ser	cga Arg	gcc Ala 290	aag Lys	cct Pro	978
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														tgc Cys		1122
					gtg Val							taat	agt	ttg		1168
gaag	gccg	cac a	agctt	gaco	et go	gaago	cacco	cto	geced	cctt	ttca	aggga	att ·	tttat	ctcga	1228
ggco	ctttc	gga ç	ggago	cagto	gg to	gggg	gtago	tgt:	caco	ctcc	aggt	atga	att (gaggg	gaggaa	1288
teg	ggtag	gaa a	actct	ccac	ga co	cato	geete	caa	atggo	cagg	atgo	ctgcc	ctt ·	tccca	acctga	1348
gago	ggad	ccc t	gtco	catgt	ig ca	gcct	cato	c aga	agcct	cac	ccto	ggag	gga :	tgeeg	gtggcg	1408
tctc	ectec	cca ç	ggago	ccaga	at ca	igtgo	cgagt	gto	gacto	jaaa	atgo	ectca	atc a	actta	agcac	1468
caaa	igcca	ıgt ç	gatca	gcaç	ga ta	ttct	gtto	cto	gtgtc	cttc	tgtt	tttt	tc	tggtg	gaatcg	1528

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<213> Homo sapiens

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Val Arg Tyr Leu Gly Tyr Leu Ala Arg Ile Asn Leu Leu Val Ala Ile $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Cys Leu Gly Leu Tyr Val Arg Trp Glu Lys Thr Ala As
n Ser Leu Ile 50 $\,\,^{<}$

Leu Val Ile Phe Ile Leu Gly Leu Phe Val Leu Gly Ile Ala Ser Ile 65 70 75 80

Leu Tyr Tyr Phe Ser Met Glu Ala Ala Ser Leu Ser Leu Ser Asn 85 90 95

Leu Trp Phe Gly Phe Leu Leu Gly Leu Leu Cys Phe Leu Asp Asn Ser 100 105 110

Ser Phe Lys Asn Asp Val Lys Glu Glu Ser Thr Lys Tyr Leu Leu 120 Thr Ser Ile Val Leu Arg Ile Leu Cys Ser Leu Val Glu Arg Ile Ser Gly Tyr Val Arg His Arg Pro Thr Leu Leu Thr Thr Val Glu Phe Leu 155 Glu Leu Val Gly Phe Ala Ile Ala Ser Thr Thr Met Leu Val Glu Lys 165 170 Ser Leu Ser Val Ile Leu Leu Val Val Ala Leu Ala Met Leu Ile Ile 185 Asp Leu Arg Met Lys Ser Phe Leu Ala Ile Pro Asn Leu Val Ile Phe 200 Ala Val Leu Leu Phe Phe Ser Ser Leu Glu Thr Pro Lys Asn Pro Ile 215 220 Ala Phe Ala Cys Phe Phe Ile Cys Leu Ile Thr Asp Pro Phe Leu Asp Ile Tyr Phe Ser Gly Leu Ser Val Thr Glu Arg Trp Lys Pro Phe Leu 245 250 Tyr Arg Gly Arg Ile Cys Arg Arg Leu Ser Val Val Phe Ala Gly Met Ile Glu Leu Thr Phe Phe Ile Leu Ser Ala Phe Lys Leu Arg Asp Thr His Leu Trp Tyr Phe Val Ile Pro Gly Phe Ser Ile Phe Gly Ile Phe 295 Trp Met Ile Cys His Ile Ile Phe Leu Leu Thr Leu Trp Gly Phe His 310 315 Thr Lys Leu Asn Asp Cys His Lys Val Tyr Phe Thr His Arg Thr Asp 330 Tyr Asn Ser Leu Asp Arg Ile Met Ala Ser Lys Gly Met Arg His Phe Cys Leu Ile Ser Glu Gln Leu Val Phe Phe Ser Leu Leu Ala Thr Ala 360 Ile Leu Gly Ala Val Ser Trp Gln Pro Thr Asn Gly Ile Phe Leu Ser Met 'Phe Leu Ile Val Leu Pro Leu Glu Ser Met Ala His Gly Leu Phe 390 395 His Glu Leu Gly Asn Cys Leu Gly Gly Thr Ser Val Gly Tyr Ala Ile 410

425 Pro Pro Glu His Val Gln Glu Leu Asn Leu Arg Ser Thr Gly Met Leu Asn Ala Ile Gln Arg Phe Phe Ala Tyr His Met Ile Glu Thr Tyr Gly 455 Cys Asp Tyr Ser Thr Ser Gly Leu Ser Phe Asp Thr Leu His Ser Lys 470 Leu Lys Ala Phe Leu Glu Leu Arg Thr Val Asp Gly Pro Arg His Asp 490 Thr Tyr Ile Leu Tyr Tyr Ser Gly His Thr His Gly Thr Gly Glu Trp 505 Ala Leu Ala Gly Gly Asp Thr Leu Arg Leu Asp Thr Leu Ile Glu Trp 520 Trp Arg Glu Lys Asn Gly Ser Phe Cys Ser Arg Leu Ile Ile Val Leu 535 Asp Ser Glu Asn Ser Thr Pro Trp Val Lys Glu Val Arg Lys Ile Asn Asp Gln Tyr Ile Ala Val Gln Gly Ala Glu Leu Ile Lys Thr Val Asp Ile Glu Glu Ala Asp Pro Pro Gln Leu Gly Asp Phe Thr Lys Asp Trp 580 Val Glu Tyr Asn Cys Asn Ser Ser Asn Asn Ile Cys Trp Thr Glu Lys 600 Gly Arg Thr Val Lys Ala Val Tyr Gly Val Ser Lys Arg Trp Ser Asp Tyr Thr Leu His Leu Pro Thr Gly Ser Asp Val Ala Lys His Trp Met Leu His Phe Pro Arg Ile Thr Tyr Pro Leu Val His Leu Ala Asn Trp 645 Leu Cys Gly Leu Asn Leu Phe Trp Ile Cys Lys Thr Cys Phe Arg Cys 665

Leu Lys Arg Leu Lys Met Ser Trp Phe Leu Pro Thr Val Leu Asp Thr
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Gly Gln Gly Phe Lys Leu Val Lys Ser

Val Ile Pro Thr Asn Phe Cys Ser Pro Asp Gly Gln Pro Thr Leu Leu

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tat tta gca atg aca aga ctg gaa gaa gta aat aga gaa gtg aac atg
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Tyr Leu Ala Met Thr Arg Leu Glu Glu Val Asn Arg Glu Val Asn Met
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His Ser Ser Val Arg Tyr Leu Gly Tyr Leu Ala Arg Ile Asn Leu Leu
gtt gct ata tgc tta ggt cta tac gta aga tgg gaa aaa aca gca aat
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Ser Leu Ile Leu Val Ile Phe Ile Leu Gly Leu Phe Val Leu Gly Ile
gcc agc ata ctc tat tac tat ttt tca atg gaa gca gca agt tta agt
                                                                   650
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ctc tcc aat ctt tgg ttt gga ttc ttg ctt ggc ctc cta tgt ttt ctt
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Leu Ser Asn Leu Trp Phe Gly Phe Leu Leu Gly Leu Leu Cys Phe Leu
                        1.00
gat aat tca tcc ttt aaa aat gat gta aaa gaa gaa tca acc aaa tat
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Asp Asn Ser Ser Phe Lys Asn Asp Val Lys Glu Glu Ser Thr Lys Tyr
                    115
                                        120
ttg ctt cta aca tcc ata gtg tta agg ata ttg tgc tct ctg gtg gag
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Leu Leu Thr Ser Ile Val Leu Arg Ile Leu Cys Ser Leu Val Glu
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										tta Leu 200						986
										tca Ser	_	_				1034
										tgc Cys						1082
										gta Val						1130
		_		_		-		-	_	aga Arg			_			1178
-		_								ctt Leu 280		-				1226
_	_							_		cct Pro						1274
										ttt Phe					22	1322
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			_	-				_	_	gtg Val 360			-			1466
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ggg Gly	ctc Leu	ttc Phe 400	cat His	gaa Glu	ttg Leu	ggt Gly	aac Asn 405	tgt Cys	tta Leu	gga Gly	gga Gly	aca Thr 410	tct Ser	gtt Val	gga Gly	1610
tat Tyr	gct Ala 415	att Ile	gtg Val	att Ile	ccc Pro	acc Thr 420	aac Asn	ttc Phe	tgc Cys	agt Ser	cct Pro 425	gat Asp	ggt Gly	cag Gln	cca Pro	1658
aca Thr 430	ctg Leu	ctt Leu	ccc Pro	cca Pro	gaa Glu 435	cat His	gta Val	cag Gln	gag Glu	tta Leu 440	aat Asn	ttg Leu	agg Arg	tct Ser	act Thr 445	1706
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acc Thr	tat Tyr	gga Gly	tgt Cys 465	gac Asp	tat Tyr	tcc Ser	aca Thr	agt Ser 470	gga Gly	ctg Leu	tca Ser	ttt	gat Asp 475	act Thr	ctg Leu	1802
cat His	tcc Ser	aaa Lys 480	cta Leu	aaa Lys	gct Ala	ttc Phe	ctc Leu 485	gaa Glu	ctt Leu	cgg Arg	aca Thr	gtg Val 490	gat Asp	gga Gly	ccc Pro	1850
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aca. Thr	gta Val 575	gat Asp	att Ile	gaa Glu	gaa Glu	gct Ala 580	gac Asp	ccg Pro	cca Pro	cag Gln	cta Leu 585	ggt Gly	gac Asp	ttt Phe	aca Thr	2138
aaa Lys	gac Asp	tgg Trp	gta Val	gaa Glu	tat Tyr	aac Asn	tgc Cys	aac Asn	tcc Ser	agt Ser	aat Asn	aac Asn	atc Ile	tgc Cys	tgg Trp	2186

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tgg agt gac tac a Trp Ser Asp Tyr T 625	ct ctg cat ttg hr Leu His Leu	cca acg gga Pro Thr Gly 630	agc gat gtg gcc Ser Asp Val Ala 635	aag 2282 Lys
cac tgg atg tta c His Trp Met Leu H 640				
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ttt agg tgc ttg a Phe Arg Cys Leu L 670	aa aga tta aaa ys Arg Leu Lys 675	atg agt tgg Met Ser Trp 680	ttt ctt cct act Phe Leu Pro Thr	gtg 2426 Val 685
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<211> 611

<212> PRT

<213> Homo sapiens

<400> 79

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- Ile Leu Cys Ser Leu Val Glu Arg Ile Ser Gly Tyr Val Arg His Arg 50 55 60
- Pro Thr Leu Leu Thr Thr Val Glu Phe Leu Glu Leu Val Gly Phe Ala 65 70 75 80
- Ile Ala Ser Thr Thr Met Leu Val Glu Lys Ser Leu Ser Val Ile Leu 85 90 95
- Leu Val Val Ala Leu Ala Met Leu Ile Ile Asp Leu Arg Met Lys Ser 100 105 110
- Phe Leu Ala Ile Pro Asn Leu Val Ile Phe Ala Val Leu Leu Phe Phe 115 120 125
- Ser Ser Leu Glu Thr Pro Lys Asn Pro Ile Ala Phe Ala Cys Phe Phe 130 135 140
- Ile Cys Leu Ile Thr Asp Pro Phe Leu Asp Ile Tyr Phe Ser Gly Leu 145 150 155 160
- Ser Val Thr Glu Arg Trp Lys Pro Phe Leu Tyr Arg Gly Arg Ile Cys 165 170 175
- Arg Arg Leu Ser Val Val Phe Ala Gly Met Ile Glu Leu Thr Phe Phe 180 185 190
- Ile Leu Ser Ala Phe Lys Leu Arg Asp Thr His Leu Trp Tyr Phe Val 195 200 205
- Ile Pro Gly Phe Ser Ile Phe Gly Ile Phe Trp Met Ile Cys His Ile 210 215 220
- Ile Phe Leu Leu Thr Leu Trp Gly Phe His Thr Lys Leu Asn Asp Cys 235 230 235
- His Lys Val Tyr Phe Thr His Arg Thr Asp Tyr Asn Ser Leu Asp Arg 245 250 255
- Ile Met Ala Ser Lys Gly Met Arg His Phe Cys Leu Ile Ser Glu Gln 260 265 270
- Leu Val Phe Phe Ser Leu Leu Ala Thr Ala Ile Leu Gly Ala Val Ser 275 280 285
- Trp Gln Pro Thr Asn Gly Ile Phe Leu Ser Met Phe Leu Ile Val Leu 290 295 300
- Pro Leu Glu Ser Met Ala His Gly Leu Phe His Glu Leu Gly Asn Cys 305 310 315 320
- Leu Gly Gly Thr Ser Val Gly Tyr Ala Ile Val Ile Pro Thr Asn Phe 325 330 335

Cys Ser Pro Asp Gly Gln Pro Thr Leu Leu Pro Pro Glu His Val Gln 340 345 350

Glu Leu Asn Leu Arg Ser Thr Gly Met Leu Asn Ala Ile Gln Arg Phe 355 360 365

Phe Ala Tyr His Met Ile Glu Thr Tyr Gly Cys Asp Tyr Ser Thr Ser 370 375 380

Gly Leu Ser Phe Asp Thr Leu His Ser Lys Leu Lys Ala Phe Leu Glu 385 390 395 400

Leu Arg Thr Val Asp Gly Pro Arg His Asp Thr Tyr Ile Leu Tyr Tyr 405 410 415

Ser Gly His Thr His Gly Thr Gly Glu Trp Ala Leu Ala Gly Gly Asp 420 425 430

Thr Leu Arg Leu Asp Thr Leu Ile Glu Trp Trp Arg Glu Lys Asn Gly 435 440 445°

Ser Phe Cys Ser Arg Leu Ile Ile Val Leu Asp Ser Glu Asn Ser Thr 450 455 460

Pro Trp Val Lys Glu Val Arg Lys Ile Asn Asp Gln Tyr Ile Ala Val 465 470 475 480

Gln Gly Ala Glu Leu Ile Lys Thr Val Asp Ile Glu Glu Ala Asp Pro \$485\$

Pro Gln Leu Gly Asp Phe Thr Lys Asp Trp Val Glu Tyr Asn Cys Asn 500 505 510

Ser Ser Asn Asn Ile Cys Trp Thr Glu Lys Gly Arg Thr Val Lys Ala 515 520 525

Val Tyr Gly Val Ser Lys Arg Trp Ser Asp Tyr Thr Leu His Leu Pro 530 535 540

Thr Gly Ser Asp Val Ala Lys His Trp Met Leu His Phe Pro Arg Ile 545 550 555 560

Thr Tyr Pro Leu Val His Leu Ala Asn Trp Leu Cys Gly Leu Asn Leu 565 570 575

Phe Trp Ile Cys Lys Thr Cys Phe Arg Cys Leu Lys Arg Leu Lys Met
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Ser Asn Leu Trp Phe Gly Phe Leu Leu Gly Leu Leu Cys Phe Leu Asp
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Phe Leu Glu Leu Val Gly Phe Ala Ile Ala Ser Thr Thr Met Leu Val
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                             80
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Glu Lys Ser Leu Ser Val Ile Leu Leu Val Val Ala Leu Ala Met Leu
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att att gat ctg aga atg aaa tot tto tta got att cca aac tta gtt 🐇
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							ctt Leu 160									1132
							tgc Cys									1180
							ttt Phe									1228
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			gac Asp 380													1804
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gaa Glu	tgg Trp	tģg Trp	aga Arg	gaa Glu 445	aag Lys	aat Asn	ggt Gly	tcc Ser	ttt Phe 450	tgt Cys	tcc Ser	cgg Arg	ctt Leu	att Ile 455	atc Ile	1996
gta Val	tta Leu	gac Asp	agc Ser 460	gaa Glu	aat Asn	tca Ser	acc Thr	cct Pro 465	tgg Trp	gtg Val	aaa Lys	gaa Glu	gtg Val 470	agg Arg	aaa Lys	2044
att Ile	aat Asn	gac Asp 475	cag Gln	tat Tyr	att Ile	gca Ala	gtg Val 480	caa Gln	gga Gly	gca Ala	gag Glu	ttg Leu 485	ata Ile	aaa Lys	aca Thr	2092
gta Val	gat Asp 490	att Ile	gaa Glu	gaa Glu	gct Ala	gac Asp 495	ccg Pro	cca Pro	cag Gln	cta Leu	ggt Gly 500	gac Asp	ttt Phe	aca Thr	aaa Lys	2140
gac Asp 505	tgg Trp	gta Val	gaa Glu	tat Tyr	aac Asn 510	tgc Cys	aac Asn	tcc Ser	agt Ser	aat Asn 515	aac Asn	atc Ile	tgc Cys	tgg Trp	act Thr 520	2188
gaa Glu	aag Lys	gga Gly	cgc Arg	aca Thr 525	gtg Val	aaa Lys	gca Ala	gta Val	tat Tyr 530	ggt Gly	gtg Val	tca Ser	aaa Lys	cgg Arg 535	tgg Trp	2236
			act Thr 540	Leu												2284
tgg Trp	atg Met	tta Leu 555	cac His	ttt Phe	cct Pro	cgt Arg	att Ile 560	aca Thr	tat Tyr	ccc Pro	cta Leu	gtg Val 565	cat His	ttg Leu.	gca Ala	2332

aat tgg tta tgc ggt ctg aac ctt ttt tgg atc tgc aaa act tgt ttt 2380 Asn Trp Leu Cys Gly Leu Asn Leu Phe Trp Ile Cys Lys Thr Cys Phe 575 agg tgc ttg aaa aga tta aaa atg agt tgg ttt ctt cct act gtg ctg 2428 Arg Cys Leu Lys Arg Leu Lys Met Ser Trp Phe Leu Pro Thr Val Leu 590 595 gac aca gga caa ggc ttc aaa ctt gtc aaa tct taatttggac cccaaagcgg 2481 Asp Thr Gly Gln Gly Phe Lys Leu Val Lys Ser 605 gatattaata agcactcata ctaccaatta tcactaactt gccatttttt gtatgctgta 2541 tttttatttg tggaaaatac ettgetaett etgtagetge teteaetttg tetttetta 2601 agtaattatg gtatatataa ggcgttggga aaaaacattt tataatgaaa gtatgtaggg 2661 agtcaaatgc ttactgtaaa tgcataagag acgttaaaaa taacactgca ctttcaggaa 2721 tgtttgctta tggtcctgat tagaaagaaa cagttgtcta tgctctgcaa tggtcaatga 2781 tgaattacta atgccttatt ttctaggcat ataataatag tttagagaat gtagaccaga 2841 taaatttgtt tactgtttta agaaaactac cagtttactt acagaagatt cttttttcca 2901 aacagtaggt ttcatccaag accatttgaa gaactgcaaa ctctttctct tagaaaagaa 2961 agagggcage ctaaaataaa egcaaaattt gettataete cateae 3007

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<212> PRT

<213> Homo sapiens

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Ser Ile Ser Phe Asn Arg Glu Lys Leu Pro Ser Ser Glu Val Val Lys 35 40 45

Phe Gly Arg Asn Ser Asn Ile Cys His Tyr Thr Phe Gln Asp Lys Gln 50 55 60

Val Ser Arg Val Gln Phe Ser Leu Gln Leu Phe Lys Lys Phe Asn Ser 65 70 75 80

Ser Val Leu Ser Phe Glu Ile Lys Asn Met Ser Lys Lys Thr Asn Leu $85 \hspace{1cm} 90 \hspace{1cm} 95$

Ile Val Asp Ser Arg Glu Leu Gly Tyr Leu Asn Lys Met Asp Leu Pro

100 105 110	
Tyr Arg Cys Met Val Arg Phe Gly Glu Tyr Gln Phe Leu Met Glu Lys , 115 120 125	
Glu Asp Gly Glu Ser Leu Glu Phe Phe Glu Thr Gln Phe Ile Leu Ser 130 135 140	
Pro Arg Ser Leu Leu Gln Glu Asn Asn Trp Pro Pro His Arg Pro Ile 145 150 155 160	
Pro Glu Tyr Gly Thr Tyr Ser Leu Cys Ser Ser Gln Ser Ser Pro 165 170 175	
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ccgcaccccg gggctcacac ttacccgcgc ggaggagcag cggccgggtg tccaccccca :	180
teetgegeee agteteeteg atteceeteg etetgageeg ggagageega acagetgaag	240
agagtteact gacteceeag ecceaggtgg geettgtgea cate atg acc agt ttt 2 Met Thr Ser Phe 1	296
gaa gat gct gac aca gaa gag aca gta act tgt ctc cag atg acg gtt Glu Asp Ala Asp Thr Glu Glu Thr Val Thr Cys Leu Gln Met Thr Val 5	344
tac cat cct ggc cag ttg cag tgt gga ata ttt cag tca ata agt ttt Tyr His Pro Gly Gln Leu Gln Cys Gly Ile Phe Gln Ser Ile Ser Phe 25 30 35	392
aac aga gag aaa ctc cct tcc agc gaa gtg gtg aaa ttt ggc cga aat 48 Asn Arg Glu Lys Leu Pro Ser Ser Glu Val Val Lys Phe Gly Arg Asn 40 45 50	140
tcc aac atc tgt cat tat act ttt cag gac aaa cag gtt tcc cga gtt Ser Asn Ile Cys His Tyr Thr Phe Gln Asp Lys Gln Val Ser Arg Val	188
cag ttt tct ctg cag ctg ttt aaa aaa ttc aac agc tca gtt ctc tcc	36

Gln	Phe 70	Ser	Leu	Gln	Leu	Phe 75	Lys	Lys	Phe	Asn	Ser 80	Ser	Val	Leu	Ser	
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aga Arg	gag Glu	ctg Leu	ggc Gly	tac Tyr 105	cta Leu	aat Asn	aaa Lys	atg Met	gac Asp 110	ctg Leu	cca Pro	tac Tyr	agg Arg	tgc Cys 115	atg Met	632
gtc Val	aga Arg	ttc Phe	gga Gly 120	gag Glu	tat Tyr	cag Gln	ttt Phe	ctg Leu 125	atg Met	gag Glu	aag Lys	gaa Glu	gat Asp 130	ggc Gly	gag Glu	680
tca Ser	ttg Leu	gaa Glu 135	ttt Phe	ttt Phe	gag Glu	act Thr	caa Gln 140	ttt Phe	att Ile	tta Leu	tct Ser	cca Pro 145	aga Arg	tca Ser	ctc Leu	728
ttg Leu	caa Gln 150	Glu	aac Asn	aac Asn	tgg Trp	cca Pro 155	cca Pro	cac His	agg Arg	ccc Pro	ata Ile 160	ccg Pro	gag Glu	tat Tyr	ggc Gly	776
act Thr 165	tat Tyr	tcg Ser	ctc Leu	tgc Cys	tcc Ser 170	tcc Ser	caa Gln	agc Ser	agt Ser	tct Ser 175	ccg Pro	aca Thr	gaa Glu	atg Met	gat Asp 180	824
		gag Glu			acac	aga .	aagt	ctaa	ga g	gaga	aata	t ga	tgga	tgaa		876
gag	ctct	gta	gatg	ctgt	at a	gaca	ctaa	a ta	agag	ttga	tta	gggt	agt	atat	tatagt	936
cat	ctgt	tat	gctg	tgaa	at t	tgga	attc	a gt	atta	tcat	ttt	gaag	tct	gtaa	attgtg	996
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ctt	atta	atc	tgtg	agga	tt a	atat	acaa	a aa	gtat	cctt	tga	gatg	aag	tcgt	gttctc	1116
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ttt	gtaa	.gga	acca	tctc	tt g	gttt	ggtc	a ca	tcag	ttca	caa	cago	cat	ttgt	tttcaa	1236
ggt	caag	gct	ccag	gcag	ıgt t	gtta	ctgg	t gt	ttgc	agco	tgt:	cagt	act	tgca	gtactg	1296
gaa	tagg	ttc	tagg	ıctag	ıtg t	ctgc	gcgt	c ac	tgtg	gttt	tag	catg	ıgga	ggac	ttattt	1356
gag	aaat	act	acct	tact	tt t	ctat	gatt	t ct	tttt	acag	ı agt	tata	ıgtg	tgtt	tactcc	1416
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aga	acca	ctg	tgtt	aagt	tt a	ggat	tatt	t ac	ttac	caaa	ı tta	ıgaaç	ıttt	gact	tttatg	1536
tgt	tata	cac	aato	cttaa	aa t	ttca	cgaa	it to	acct	tttt	aat	agta	tcc	atgt	acataa	1596
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<211> 392
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<212> PRT

<213> Homo sapiens

<400> 83

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Phe Met Val Pro Gly Tyr Leu Leu Val Gln Tyr Phe Arg Arg Lys Asn 50 55 60

Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys 65 70 75 80

Val Phe Gly Asn Glu Pro Lys Ala Ser Asp Glu Val Pro Leu Ala Pro 85 90 95

Arg Thr Glu Ala Ala Glu Thr Thr Pro Met Trp Gln Ala Leu Lys Leu 100 . 105 110

Leu Phe Cys Ala Thr Gly Leu Gln Val Ser Tyr Leu Thr Trp Gly Val 115 120 125

Leu Gln Glu Arg Val Met Thr Arg Ser Tyr Gly Ala Thr Ala Thr Ser 130 135 140

Pro Gly Glu Arg Phe Thr Asp Ser Gln Phe Leu Val Leu Met Asn Arg 145 150 155 160

Val Leu Ala Leu Ile Val Ala Gly Leu Ser Cys Val Leu Cys Lys Gln 165 170 175

Pro Arg His Gly Ala Pro Met Tyr Arg Tyr Ser Phe Ala Ser Leu Ser 180 185 190

Asn Val Leu Ser Ser Trp Cys Gln Tyr Glu Ala Leu Lys Phe Val Ser 195 200 205

Phe Pro Thr Gln Val Leu Ala Lys Ala Ser Lys Val Ile Pro Val Met 210 215 220

Leu Met Gly Lys Leu Val Ser Arg Arg Ser Tyr Glu His Trp Glu Tyr 225 230 235 240

Leu Thr Ala Thr Leu Ile Ser Ile Gly Val Ser Met Phe Leu Leu Ser 245 250 255

Ser Gly Pro Glu Pro Arg Ser Ser Pro Ala Thr Thr Leu Ser Gly Leu 260 265 270

Phe	Thr	Val 275	GLY	Ser	Leu	Leu	G1u 280	GIn	GLY	Ala	Leu	ьеи 285	GLu	GLY	Thr	
Arg	Phe 290	Met	Gly	Arg	His	Ser 295	Glu	Phe	Ala	Ala	His 300	Ala	Leu	Leu	Leu	
Ser 305	Ile	Cys	Ser	Ala	Cys 310	Gly	Gln	Leu	Phe	Ile 315	Phe	Tyr	Thr	Ile	Gly 320	
Gln	Phe	Gly	Ala	Ala 325	Val	Phe	Thr	Ile	Ile 330	Met	Thr	Leu	Arg	Gln 335	Ala	
Phe	Ala	Ile	Leu 340	Leu	Ser	Cys	Leu	Leu 345	Tyr	Gly	His	Thr	Val 350	Thr	Val	
Val	Gly	Gly 355	Leu	Gly	Val	Ala	Val 360	Val	Phe	Ala	Ala	Leu 365	Leu	Leu	Arg	
Val	Tyr 370	Ala	Arg	Gly	Arg	Leu 375	Lys	Gln	Arg	Gly	Lys 380	Lys	Ala	Val	Pro	
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cgc	gcgg	ccc (gggg	actc	gc at	ttcc	ccgg	t tc	cccc	tcca	ccc	cacg	cgg (cctg	gacc	118
														ccc Pro 15		166
cta Leu	ggg Gly	gca Ala	ggt Gly 20	ggg	gag Glu	act Thr	ccc Pro	gaa Glu 25	gcc Ala	cct Pro	ccg Pro	gag Glu	tca Ser 30	tgg Trp	acc Thr	214
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ttt Phe	atg Met 50	gta Val	cct Pro	ggc	tac Tyr	ctc Leu 55	ctg Leu	gtg Val	cag Gln	tac Tyr	ttc Phe 60	agg Arg	cgg Arg	aag Lys	aac Asn	310
														gct Ala		358

65				**	70					75					80	
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cga Arg	aca Thr	gag Glu	gcg Ala 100	gca Ala	gag Glu	acc Thr	acc Thr	ccg Pro 105	atg Met	tgg Trp	cag Gln	gcc Ala	ctg Leu 110	aag Lys	ctg Leu	454
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ctg Leu	cag Gln 130	gaa Glu	aga Arg	gtg Val	atg Met	acc Thr 135	cgc Arg	agc Ser	tat Tyr	ggg Gly	gcc Ala 140	aca Thr	gcc Ala	aca Thr	tca Ser	550
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gtg Val	ctg Leu	gca Ala	ctg Leu	att Ile 165	gtg Val	gct Ala	ggc Gly	ctc Leu	tcc Ser 170	tgt Cys	gtt Val	ctc Leu	tgc Cys	aag Lys 175	cag Gln	646
ccc Pro	cgg Arg	cat His	ggg Gly 180	gca Ala	ccc Pro	atg Met	tac Tyr	cgg Arg 185	tac Tyr	tcc Ser	ttt Phe	gcc Ala	agc Ser 190	ctg Leu	tcc Ser	694
aat Asn	gtg Val	ctt Leu 195	agc Ser	agc Ser	tgg Trp	tgc Cys	caa Gln 200	tac Tyr	gaa Glu	gct Ala	ctt Leu	aag Lys 205	ttc Phe	gtc Val	agc Ser	7.42
ttc Phe	ccc Pro 210	acc Thr	cag Gln	gtg Val	ctg Leu	gcc Ala 215	aag Lys	gcc Ala	tct Ser	aag Lys	gtg Val 220	atc Ile	cct Pro	gtc Val	atg Met	790
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ctg Leu	aca Thr	gcc Ala	acc Thr	ctc Leu 245	atc Ile	tcc Ser	att Ile	gly	gtc Val 250	agc Ser	atg Met	ttt Phe	ctg Leu	cta Leu 255	tcc Ser	886
agc Ser	gga Gly	cca Pro	gag Glu 260	Pro	cgc Arg	agc Ser	tcc Ser	cca Pro 265	Ala	acc Thr	aca Thr	ctc Leu	tca Ser 270	ggc Gly	ctc Leu	934
ttc Phe	aca Thr	gtg Val 275	Gly	tca Ser	ctg Leu	cta Leu	gaa Glu 280	cag Gln	ggg Gly	gcc Ala	cta Leu	ctg Leu 285	gag Glu	gga Gly	acc Thr	982
cgc Arg	ttc Phe 290	Met	Gly	cga Arg	cac His	agt Ser 295	Glu	ttt Phe	gct Ala	gcc Ala	cat His 300	Ala	ctg Leu	cta Leu	ctc Leu	1030

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tcc atc tgc tcc gca tgt ggc cag ctc ttc atc ttt tac acc att ggg
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Ser Ile Cys Ser Ala Cys Gly Gln Leu Phe Ile Phe Tyr Thr Ile Gly
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cag ttt ggg get gee gte tte ace ate ate atg ace ete ege cag gee
                                                                1126
Gln Phe Gly Ala Ala Val Phe Thr Ile Ile Met Thr Leu Arg Gln Ala
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                                   330
ttt gcc atc ctt ctt tcc tgc ctt ctc tat ggc cac act gtc act gtg
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Phe Ala Ile Leu Leu Ser Cys Leu Leu Tyr Gly His Thr Val Thr Val
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                               345
                                                  350
1222
Val Gly Gly Leu Gly Val Ala Val Val Phe Ala Ala Leu Leu Leu Arg
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                           360
                                              365
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                                                                1270
Val Tyr Ala Arg Gly Arg Leu Lys Gln Arg Gly Lys Lys Ala Val Pro
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gtt gag tet eet gtg eag aag gtt tgagggtgga aagggeetga ggggtgaagt
Val Glu Ser Pro Val Gln Lys Val
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gaaataggac cctcccacca tccccttctg ctgtaacctc tgagggagct ggctgaaagg 1384
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<212> PRT

<213> Homo sapiens

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Phe Met Val Pro Gly Tyr Leu Leu Val Gln Tyr Phe Arg Arg Lys Asn Tyr Leu Glu Thr Gly Arg Gly Leu Cys Phe Pro Leu Val Lys Ala Cys Val Phe Gly Asn Glu Pro Lys Ala Ser Asp Glu Val Pro Leu Ala Pro Arg Thr Glu Ala Ala Glu Thr Thr Pro Met Trp Gln Ala Leu Lys Leu 105 Leu Phe Cys Ala Thr Gly Leu Gln Val Ser Tyr Leu Thr Trp Gly Val 120 Leu Gln Glu Arg Val Met Thr Arg Ser Tyr Gly Ala Thr Ala Thr Ser 135 Pro Gly Glu Arg Phe Thr Asp Ser Gln Phe Leu Val Leu Met Asn Arg Val Leu Ala Leu Ile Val Ala Gly Leu Ser Cys Val Leu Cys Lys Gln Pro Arg His Gly Ala Pro Met Tyr Arg Tyr Ser Phe Ala Ser Leu Ser Asn Val Leu Ser Ser Trp Cys Gln Tyr Glu Ala Leu Lys Phe Val Ser 195 Phe Pro Thr Gln Val Leu Ala Lys Ala Ser Lys Val Ile Pro Val Met 215 Leu Met Gly Lys Leu Val Ser Arg Arg Ser Tyr Glu His Trp Glu Tyr 230 235

Gln Leu Trp Phe Phe Arg Phe Val Val Asn Ala Ala Gly Tyr Ala Ser

Gln Asp Ala Leu Phe Ala Tyr Lys Met Ser Ser Val Gln Met Met Phe 290 295 300

Leu Thr Ala Thr Leu Ile Ser Ile Gly Val Ser Met Phe Leu Leu Ser

Ser Gly Pro Glu Pro Arg Ser Ser Pro Ala Thr Thr Leu Ser Gly Leu

Ile Leu Leu Ala Gly Tyr Ile Ala Phe Asp Ser Phe Thr Ser Asn Trp 275 280 285

- Gly Val Asn Phe Phe Ser Cys Leu Phe Thr Val Gly Ser Leu Leu Glu 305 310 315 320
- Gln Gly Ala Leu Leu Glu Gly Thr Arg Phe Met Gly Arg His Ser Glu 325 330 335

Phe	Ala	Ala	His 340	Ala	Leu	Leu	Leu	Ser 345	Ile	Суѕ	Ser	Ala	Cys 350	Gly	Gln	
Leu	Phe	Ile 355	Phe	Tyr	Thr	Ile	Gly 360	Gln	Phe	Gly	Ala	Ala 365	Val	Phe	Thr	
Ile	Ile 370	Met	Thr	Leu	Arg	Gln 375	Ala	Phe	Ala	Ile	Leu 380	Leu	Ser	Cys	Leu	
Leu 385	Tyr	Gly	His	Thr	Val 390	Thr	Val	Val	Gly	Gly 395	Leu	Gly	Val	Ala	Val 400	
Val	Phe	Ala	Ala	Leu 405	Leu	Leu	Arg	Val	Tyr 410	Ala	Arg	Gly	Arg	Leu 415	Lys	
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cgc	gcgg	ccc (gggg	actc	gc a	ttcc	ccggt	t to	cccc.	tcca	ccc	cacg	cgg	cctg	gacc	118
atg Met 1	Āsp	gcc Ala	aga Arg	tgg Trp 5	tgg Trp	gca Ala	gtg Val	gtg Val	gtg Val 10	ctg Leu	gct Ala	gcg Ala	ttc Phe	ccc Pro 15	tcc Ser	166
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cag Gln	cta Leu	tgg Trp 35	ttc Phe	ttc Phe	cga Arg	ttt Phe	gtg Val 40	gtg Val	aat Asn	gct Ala	gct Ala	ggc Gly 45	tat Tyr	gcc Ala	agc Ser	262
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tac Tyr 65	Leu	gag Glu	acc Thr	ggt Gly	agg Arg 70	ggc Gly	ctc Leu	tgc Cys	ttt Phe	ccc Pro 75	Leu	gtg Val	aaa Lys	gct Ala	tgt Cys 80	358
gtg																

														aag Lys		454
		_	_				-				_			ggt Gly		502
_	_	_			_		-	-			_		-	aca Thr		550
														aac Asn		598
gtg Val	ctg Leu	gca Ala	ctg Leu	att Ile 165	gtg Val	gct Ala	ggc Gly	ctc Leu	tcc Ser 170	tgt Cys	gtt Val	ctc Leu	tgc Cys	aag Lys 175	cag Gln	646
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														gtc Val		742
														gtc Val		790
														gag Glu		838
														cta Leu 255		886
														ggc Gly		934
														aac Asn		982
cag Gln	gat Asp 290	gcc Ala	ctg Leu	ttt Phe	gcc Ala	tat Tyr 295	aag Lys	atg Met	tca Ser	tcg Ser	gtg Val 300	cag Gln	atg Met	atg Met	ttt Phe	1030
											Gly			cta Leu		1078

cag ggg gcc o Gln Gly Ala 1						1126
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ctc ttc atc t Leu Phe Ile : 355						1222
atc atc atg a Ile Ile Met ' 370	_		-		-	1270
ctc tat ggc Leu Tyr Gly : 385	_					1318
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çaa cgg gga Gln Arg Gly			. Glu Ser		Lys Val	1414
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caaatacctg c	atttettae e	ctggtgaga aa	agcacaag	cggtgtaggc	tccaatgctg	1834
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<212> PRT

<213> Homo sapiens

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Ser Asn Thr Glu Gly Pro Thr Gly Lys Gln Glu Gly Ala Gln Ser 50 60

Val Glu Glu Met Phe Glu Glu Glu Glu Glu Glu Val Phe Leu Lys
65 70 75 80

Phe Val Ile Leu His Ala Glu Asp Asp Thr Asp Glu Ala Leu Arg Val85 90 95

Gln Asn Leu Gln Asp Asp Phe Gly Ile Lys Pro Gly Ile Ile Phe 100 105 110

Ala Glu Met Pro Cys Gly Arg Gln His Leu Gln Asn Leu Asp Asp Ala 115 120 125

Val Asn Gly Ser Ala Trp Thr Ile Leu Leu Leu Thr Glu Asn Phe Leu 130 \$135\$ 140 ,

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Ile Ala Thr Gly Thr Val Ile Ile Leu Leu Gly Thr Phe Gly Cys Phe 65 70 75 80

Ala Thr Cys Arg Ala Ser Ala Trp Met Leu Lys Leu Tyr Ala Met Phe \$85\$ 90 95

Leu Thr Leu Val Phe Leu Val Glu Leu Val Ala Ala Ile Val Gly Phe 100 $$ 105 $$ 110

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<213> Homo sapiens

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Cys Leu Glu Gly Ile Leu Glu Gly Ser Val Arg Asn Ser Leu Trp Arg 35 40 45

Pro Ala Pro Phe Lys Cys Pro Thr Cys Arg Lys Glu Thr Ser Ala Thr Gly Ile Asn Ser Leu Gln Val Asn Tyr Ser Leu Lys Gly Ile Val Glu Lys Tyr Asn Lys Ile Lys Ile Ser Pro Lys Met Pro Val Cys Lys Gly His Leu Gly Gln Pro Leu Asn Ile Phe Cys Leu Thr Asp Met Gln Leu 105 Ile Cys Gly Ile Cys Ala Thr Arg Gly Glu His Thr Lys His Val Phe 120 Cys Ser Ile Glu Asp Ala Tyr Ala Gln Glu Arg Asp Ala Phe Glu Ser Leu Phe Gln Ser Phe Glu Thr Trp Arg Arg Gly Asp Ala Leu Ser Arg Leu Asp Thr Leu Glu Thr Ser Lys Arg Lys Ser Leu Gln Leu Leu Thr Lys Asp Ser Asp Lys Val Lys Glu Phe Phe Glu Lys Leu Gln His Thr 185 Leu Asp Gln Lys Lys Asn Glu Ile Leu Ser Asp Phe Glu Thr Met Lys Leu Ala Val Met Gln Ala Tyr Asp Pro Glu Ile Asn Lys Leu Asn Thr Ile Leu Gln Glu Gln Arg Met Ala Phe Asn Ile Ala Glu Ala Phe Lys 235 Asp Val Ser Glu Pro Ile Val Phe Leu Gln Gln Met Gln Glu Phe Arg Glu Lys Ile Lys Val Ile Lys Glu Thr Pro Leu Pro Pro Ser Asn Leu Pro Ala Ser Pro Leu Met Lys Asn Phe Asp Thr Ser Gln Trp Glu Asp Ile Lys Leu Val Asp Val Asp Lys Leu Ser Leu Pro Gln Asp Thr Gly 295 300 Thr Phe Ile Ser Lys Ile Pro Trp Ser Phe Tyr Lys Leu Phe Leu Leu 315 Ile Leu Leu Gly Leu Val Ile Val Phe Gly Pro Thr Met Phe Leu 325 330 Glu Trp Ser Leu Phe Asp Asp Leu Ala Thr Trp Lys Gly Cys Leu Ser

Asn Phe Ser Ser Tyr Leu Thr Lys Thr Ala Asp Phe Ile Glu Gln Ser Val Phe Tyr Trp Glu Gln Val Thr Asp Gly Phe Phe Ile Phe Asn Glu Arg Phe Lys Asn Phe Thr Leu Val Val Leu Asn Asn Val Ala Glu Phe 390 395 Val Cys Lys Tyr Lys Leu Leu 405 <210> 96 <211> 1409 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (181)..(1401) <400> 96 gctgtgcttg gcgcgtaccg tgcggtccct gtagttggag gacgggcggt cgcgcggcct 60 ttcccactag ccggagtagc ctctagttcg ttagtcaaaa cgtgaaaaaa aaagacctgc 120 tttgccctgg gaaatagtaa ccctgccaaa tacatcagct tgtaggagac agaggatgtg 180 atg gag ctg ctt gaa gaa gat ctc aca tgc cct att tgt tgt agt ctg 228 Met Glu Leu Leu Glu Glu Asp Leu Thr Cys Pro Ile Cys Cys Ser Leu 1 5 ttt gat gat cca cgg gtt ttg cct tgc tcc cac aac ttc tgc aaa aaa 276 Phe Asp Asp Pro Arg Val Leu Pro Cys Ser His Asn Phe Cys Lys 20 25 30 tgc tta gaa ggt atc tta gaa ggg agt gtg cgg aat tcc ttg tgg aga 324 Cys Leu Glu Gly Ile Leu Glu Gly Ser Val Arg Asn Ser Leu Trp Arg 35 40 cca gct cca ttc aag tgt cct aca tgc cgt aag gaa act tca gct act 372 Pro Ala Pro Phe Lys Cys Pro Thr Cys Arg Lys Glu Thr Ser Ala Thr 50 gga att aat agc ctg cag gtt aat tac tcc ctg aag ggt att gtg gaa 420 Gly Ile Asn Ser Leu Gln Val Asn Tyr Ser Leu Lys Gly Ile Val Glu 65 70 aag tat aac aag atc aag atc tct ccc aaa atg cca gta tgc aaa gga 468 Lys Tyr Asn Lys Ile Lys Ile Ser Pro Lys Met Pro Val Cys Lys Gly 85 cac ttg ggg cag cct ctc aac att ttc tgc ctg act gat atg cag ctg 516 His Leu Gly Gln Pro Leu Asn Ile Phe Cys Leu Thr Asp Met Gln Leu 100

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			act Thr								708
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			aat Asn								804
			gca Ala								852
			cgg Arg 230								,900
			att Ile								948
			atc Ile								996
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			att Ile 310								1140
			ctt Leu								1188

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His Ser Tyr 35		eu Cys Ile 40	-	Phe Phe	Lys Asn 45	Pro	Ser
Gln Lys Gln 50	Leu Arg G	ln Glu Thr 55	Phe Cys	Cys Pro 60	Gln Cys	Arg	Ala
Pro Phe His 65	_	er Leu Arg 70	Pro Asn	Lys Gln 75	Leu Gly	Ser	Leu 80
Ile Glu Ala	Leu Lys G	lu Thr Asp	Gln Glu 90	Met Ser	Cys Glu	Glu 95	His
Gly Glu Gln	Phe His L	eu Phe Cys	Glu Asp 105	Glu Gly	Gln Leu 110	Ile	Cys
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Val Glu Asp 130	Val Cys G	ln Gly Tyr 135	Lys Glu	Lys Leu 140	Gln Glu	Ala	Val
Thr Lys Leu 145	-	eu Glu Asp 50	Arg Cys	Thr Glu 155	Gln Lys		Ser 160

Thr Ala Met Arg Ile Thr Lys Trp Lys Glu Lys Val Gln Ile Gln Arg 170 Gln Lys Ile Arg Ser Asp Phe Lys Asn Leu Gln Cys Phe Leu His Glu 185 Glu Glu Lys Ser Tyr Leu Trp Arg Leu Glu Lys Glu Glu Gln Gln Thr Leu Ser Arg Leu Arg Asp Tyr Glu Ala Gly Leu Gly Leu Lys Ser Asn 215 Glu Leu Lys Ser His Ile Leu Glu Leu Glu Glu Lys Cys Gln Gly Ser 235 Ala Gln Lys Leu Gln Asn Val Asn Asp Thr Leu Ser Arg Ser Trp 250 Ala Val Lys Leu Glu Thr Ser Glu Ala Val Ser Leu Glu Leu His Thr 265 Met Cys Asn Val Ser Lys Leu Tyr Phe Asp Val Lys Lys Met Leu Arg 280 Ser His Gln Val Ser Val Thr Leu Asp Pro Asp Thr Ala His His Glu Leu Ile Leu Ser Glu Asp Arg Arg Gln Val Thr Arg Gly Tyr Thr Gln 315 Glu Asn Gln Asp Thr Ser Ser Arg Arg Phe Thr Ala Phe Pro Cys Val 330 Leu Gly Cys Glu Gly Phe Thr Ser Gly Arg Arg Tyr Phe Glu Val Asp Val Gly Glu Gly Thr Gly Trp Asp Leu Gly Val Cys Met Glu Asn Val Gln Arg Gly Thr Gly Met Lys Gln Glu Pro Gln Ser Gly Phe Trp Thr Leu Arg Leu Cys Lys Lys Gly Tyr Val Ala Leu Thr Ser Pro Pro 395 Thr Ser Leu His Leu His Glu Gln Pro Leu Leu Val Gly Ile Phe Leu Asp Tyr Glu Ala Gly Val Val Ser Phe Tyr Asn Gly Asn Thr Gly Cys His Ile Phe Thr Phe Pro Lys Ala Ser Phe Ser Asp Thr Leu Arg Pro Tyr Phe Gln Val Tyr Gln Tyr Ser Pro Leu Phe Leu Pro Pro Pro Gly 455 460

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Asp

Glu	Gln	Phe 100	His	Leu	Phe	Cys	Glu 105	Asp	Glu	Gly	Gln	Leu 110	Ile	Cys	Trp	
_	_			-	cca Pro	_							_		_	863
_	-	_	_	_	ggc Gly 135	Tyr	_	_	_		_	_	_			911
					gaa Glu			_	_		_	_	_			959
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				_	ttt Phe	_			_	_					_	1055
	_				tgg Trp		_			-	_		_			1103
		_		_	tat Tyr 215		-		_		_	_			_	1151
	_	_			ctg Leu	_	_		-		-	_			-	1199
			_	_	aat Asn			_		_	_		_		_	1247
	_	_	_		tca Ser		_	_		_						1295
				_	ctt Leu			_		_		_			agt Ser	1343
				_	act Thr 295	_	_		-		_			_		1391
				_	cgg Arg	_								_		1439
	_	_			tcc Ser		_			_			_	_	_	1487

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								gga Gly								1583
								cct Pro								1631
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								ctg Leu 410								1727
								tat Tyr								1775
								ttc Phe								1823
								ttg Leu								1871
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His	Ser	Tyr 35	Cys	His	Leu	Cys	Ile 40	Thr	Asp	Phe	Phe	Lys 45	Asn	Pro	Ser	
Gln	Lys 50	Gln	Leu	Arg	Gln	Glu 55	Thr	Phe	Cys	Суѕ	Pro 60	Gln	Cys	Arg	Ala	

Pro Phe His Met Asp Ser Leu Arg Pro Asn Lys Gln Leu Gly Ser Leu Ile Glu Ala Leu Lys Glu Thr Asp Gln Glu Met Ser Cys Glu Glu His Gly Glu Gln Phe His Leu Phe Cys Glu Asp Glu Gly Gln Leu Ile Cys Trp Arg Cys Glu Arg Ala Pro Gln His Lys Gly His Thr Thr Ala Leu Val Glu Asp Val Cys Gln Gly Tyr Lys Glu Lys Leu Gln Lys Ala Val Thr Lys Leu Lys Gln Leu Glu Asp Arg Cys Thr Glu Gln Lys Leu Ser 155 Thr Ala Met Arg Ile Thr Lys Trp Lys Glu Lys Val Gln Ile Gln Arg 170 Gln Lys Ile Arg Ser Asp Phe Lys Asn Leu Gln Cys Phe Leu His Glu Glu Glu Lys Ser Tyr Leu Trp Arg Leu Glu Lys Glu Glu Gln Gln Thr 200 Leu Ser Arg Leu Arg Asp Tyr Glu Ala Gly Leu Gly Leu Lys Ser Asn Glu Leu Lys Ser His Ile Leu Glu Leu Glu Glu Lys Cys Gln Gly Ser Ala Gln Lys Leu Leu Gln Asn Val Asn Asp Thr Leu Ser Arg Ser Trp 245 Ala Val Lys Leu Glu Thr Ser Glu Ala Val Ser Leu Glu Leu His Thr 2.65 Met Cys Asn Val Ser Lys Leu Tyr Phe Asp Val Lys Lys Met Leu Arg 280 Ser His Gln Val Ser Val Thr Leu Asp Pro Asp Thr Ala His His Glu Leu Ile Leu Ser Glu Asp Arg Gln Val Thr Arg Gly Tyr Thr Gln Glu Asn Gln Asp Thr Ser Ser Arg Arg Phe Thr Ala Phe Pro Cys Val 330 Leu Gly Cys Glu Gly Phe Thr Ser Gly Arg Arg Tyr Phe Glu Val Asp Val Gly Glu Gly Thr Gly Trp Asp Leu Gly Val Cys Met Glu Asn Val 360

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Thr Ser Leu His Leu His Glu Gln Pro Leu Leu Val Gly Ile Phe Leu
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Asp Tyr Glu Ala Gly Val Val Ser Phe Tyr Asn Gly Asn Thr Gly Cys
            420
                                425
His Ile Phe Thr Phe Pro Lys Ala Ser Phe Ser Asp Thr Leu Arg Pro
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ged tea acc acc age acc aag aag atg atg gag gaa ged acc tgd ted
Ala Ser Thr Thr Ser Thr Lys Lys Met Met Glu Glu Ala Thr Cys Ser
atc tgc ctg agc ctg atg acg aac cca gta agc atc aac tgt gga cac
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Ile Cys Leu Ser Leu Met Thr Asn Pro Val Ser Ile Asn Cys Gly His
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											cag Gln					671
											ctg Leu					719
-	_										tgt Cys					767
											cag Gln					815
											acc Thr 125					863
											cag Gln					911
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cag Gln	aaa Lys	ttg Leu	ctg Leu 245	cag Gln	aat Asn	gtg Val	aat Asn	gac Asp 250	Thr	ttg Leu	agc Ser	agg Arg	agt Ser 255	tgg Trp	gct Ala	1247

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												cat His				1391
												tac Tyr				1439
												ccc Pro				1487
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												tct Ser				1679
												att Ile				1727
												act Thr 430				1775
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											Pro				gac Asp 465	1871
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Ala Gln Val Pro Pro Ala Ala Pro His His His His His Ser His 50 55 60

Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr Gly Lys
65 70 75 80

Arg Tyr Cys Arg Gly Lys Val Leu Gly Lys Gly Gly Phe Ala Lys Cys 85 90 95

Tyr Glu Met Thr Asp Leu Thr Asn Asn Lys Val Tyr Ala Ala Lys Ile 100 105 110

Ile Pro His Ser Arg Val Ala Lys Pro His Gln Arg Glu Lys Ile Asp 115 120 125

Lys Glu Ile Glu Leu His Arg Ile Leu His His Lys His Val Val Gln 130 135 140

Phe Tyr His Tyr Phe Glu Asp Lys Glu Asn Ile Tyr Ile Leu Leu Glu 145 150 155 160

Tyr Cys Ser Arg Arg Ser Met Ala His Ile Leu Lys Ala Arg Lys Val 165 170 175

Leu Thr Glu Pro Glu Val Arg Tyr Tyr Leu Arg Gln Ile Val Ser Gly 180 185 190

Leu Lys Tyr Leu His Glu Glu Glu Ile Leu His Arg Asp Leu Lys Leu 195 200 205

Gly Asn Phe Phe Ile Asn Glu Ala Met Glu Leu Lys Val Gly Asp Phe 210 215 220

Gly Leu Ala Ala Arg Leu Glu Pro Leu Glu His Arg Arg Arg Thr Ile 225 230 235 240

Cys Gly Thr Pro Asn Tyr Leu Ser Pro Glu Val Leu Asn Lys Gln Gly
245 250 255

His Gly Cys Glu Ser Asp Ile Trp Ala Leu Gly Cys Val Met Tyr Thr 260 265 270

Met Leu Leu Gly Arg Pro Pro Phe Glu Thr Thr Asn Leu Lys Glu Thr 280 Tyr Arg Cys Ile Arg Glu Ala Arg Tyr Thr Met Pro Ser Ser Leu Leu 295 300 Ala Pro Ala Lys His Leu Ile Ala Ser Met Leu Ser Lys Asn Pro Glu 310 315 Asp Arg Pro Ser Leu Asp Asp Ile Ile Arg His Asp Phe Phe Leu Gln 330 Gly Phe Thr Pro Asp Arg Leu Ser Ser Ser Cys Cys His Thr Val Pro 345 Asp Phe His Leu Ser Ser Pro Ala Lys Asn Phe Phe Lys Lys Ala Ala Ala Ala Leu Phe Gly Gly Lys Lys Asp Lys Ala Arg Tyr Ile Asp Thr His Asn Arg Val Ser Lys Glu Asp Glu Asp Ile Tyr Lys Leu Arg His 395 Asp Leu Lys Lys Thr Ser Ile Thr Gln Gln Pro Ser Lys His Arg Thr Asp Glu Glu Leu Gln Pro Pro Thr Thr Val Ala Arg Ser Gly Thr Pro Ala Val Glu Asn Lys Gln Gln Ile Gly Asp Ala Ile Arg Met Ile 440 Val Arg Gly Thr Leu Gly Ser Cys Ser Ser Ser Ser Glu Cys Leu Glu Asp Ser Thr Met Gly Ser Val Ala Asp Thr Val Ala Arg Val Leu Arg Gly Cys Leu Glu Asn Met Pro Glu Ala Asp Cys Ile Pro Lys Glu Gln Leu Ser Thr Ser Phe Gln Trp Val Thr Lys Trp Val Asp Tyr Ser Asn 505 Lys Tyr Gly Phe Gly Tyr Gln Leu Ser Asp His Thr Val Gly Val Leu Phe Asn Asn Gly Ala His Met Ser Leu Leu Pro Asp Lys Lys Thr Val 535 His Tyr Tyr Ala Glu Leu Gly Gln Cys Ser Val Phe Pro Ala Thr Asp 545 Ala Pro Glu Gln Phe Ile Ser Gln Val Thr Val Leu Lys Tyr Phe Ser 570

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Phe Tyr His Asp His Thr Lys Ile Ile Cys Ser Gln Asn Glu Glu
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Tyr Leu Leu Thr Tyr Ile Asn Glu Asp Arg Ile Ser Thr Thr Phe Arg
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Thr Lys Met Cys Glu Gln Ala Leu Gly Lys Gly Cys Gly Ala Asp Ser
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Lys Lys Arg Pro Pro Gln Pro Pro Glu Glu Ser Gln Pro Pro Gln
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Ser Gln Ala Gln Val Pro Pro Ala Ala Pro His His His His His
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Ser His Ser Gly Pro Glu Ile Ser Arg Ile Ile Val Asp Pro Thr Thr
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			gaa Glu 130													553
			tac Tyr												ctc Leu .	601
			tgc Cys													649
			aca Thr										_			697
			aaa Lys													745
			aac Asn 210													793
			ctg Leu					_		_	_		_			841
			ggt Gly								_	-				889
			ggc Gly	_	-		_			_	-		_	-		937
		_	tta Leu							_						985
			agg Arg 290													1033
			cct Pro		_				_	-	_	-				1081

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										tct Ser 345						1177
										aag Lys						1225
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agg Arg	cat His 400	gat Asp	ttg Leu	aaa Lys	aag Lys	act Thr 405	tca Ser	ata Ile	act Thr	cag Gln	caa Gln 410	ccc Pro	agc Ser	aaa Lys	cac His	1369
										acc Thr 425						1417
										att						1465
										agc Ser					tgc Cys	1513
										gac Asp						1561
										gct Ala						1609
										acc Thr 505						1657
										tca Ser					ggt Gly.	1705
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														caa Gln		351
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-		_	_			-		_		-	-	_	-	ctg Leu 125		447
														ctg Leu		495
		_			_					-		_		acc Thr		543
_	cgt Arg 160	_	tago	gacto	cag d	ccaga	acgto	gg ag	ggga	geeg	g gto	gccg	cagg			592
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ttga	agget	gt t	ctg	ccaç	ga go	etege	gctga	a acc	cagco	ettt	agto	geeta	acc	attat	tcttat	101
coat	- a+ a+	-+	acat a	agat (*	*	~ - +	- a++	- ~ ~ ~ +	-+	200	·+++		~~~++	-~~~+	107

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<213> Homo sapiens

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Tyr Pro Thr Pro Pro Ala Pro Met Pro Gly Pro Thr Thr Gly Leu Val 35 40 45

Thr Gly Pro Asp Gly Lys Gly Met Asn Pro Pro Ser Tyr Tyr Thr Gln 50 55 \cdot 60

Pro Ala Pro Ile Pro Asn Asn Asn Pro Ile Thr Val Gln Thr Val Tyr 65 70 75 80

Val Gln His Pro Ile Thr Phe Leu Asp Arg Pro Val Gln Met Cys Cys
85 90 95

Pro Ser Cys Asn Lys Met Ile Val Ser Gln Leu Ser Tyr Asn Ala Gly 100 105 110

Ala Leu Thr Trp Leu Ser Cys Gly Ser Leu Cys Leu Leu Gly Cys Ile 115 120 . 125

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Leu

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	L> CI	os 70).	. (552	2)												
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				aca Thr 35												207
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				ccc Pro										_	_	303
				cac His												351
				tgc Cys		-	_			-	-	_				399
				acc Thr 115												447
				tgc Cys												495
				tac Tyr												543
aag Lvs			tag	gacto	cag o	ccaga	acgt	gg ag	ggga	gccg	g gto	gccg	cagg			592

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Val Thr Thr Asn Leu Lys Leu Arg Asn Pro Ser Asp Arg Lys Val Cys
35 40 45

Phe Lys Val Lys Thr Thr Ala Pro Arg Arg Tyr Cys Val Arg Pro Asn 50 55 60

Ser Gly Ile Ile Asp Pro Gly Ser Thr Val Thr Val Ser Val Met Leu 65 70 75 80

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Gln Pro Phe Asp Tyr Asp Pro Asn Glu Lys Ser Lys His Lys Phe Met
Val Gln Thr Ile Phe Ala Pro Pro Asn Thr Ser Asp Met Glu Ala Val
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Trp Lys Glu Ala Lys Pro Asp Glu Leu Met Asp Ser Lys Leu Arg Cys
       115
                            120
Val Phe Glu Met Pro Asn Glu Asn Asp Lys Leu Asn Asp Met Glu Pro
                        135
                                            140
Ser Lys Ala Val Pro Lèu Asn Ala Ser Lys Gln Asp Gly Pro Met Pro
                    150
                                        155
Lys Pro His Ser Val Ser Leu Asn Asp Thr Glu Thr Arg Lys Leu Met
                                    170
Glu Glu Cys Lys Arg Leu Gln Gly Glu Met Met Lys Leu Ser Glu Glu
Asn Arg His Leu Arg Asp Glu Gly Leu Arg Leu Arg Lys Val Ala His
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Ser Asp Lys Pro Gly Ser Thr Ser Thr Ala Ser Phe Arg Asp Asn Val
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Thr Ser Pro Leu Pro Ser Leu Leu Val Val Ile Ala Ala Ile Phe Ile
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Gly Phe Phe Leu Gly Lys Phe Ile Leu
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gegageetgg cetegteeta gageteggee gageegtege egeegtegte eeeegeecee 180
agteageaaa eegeegeege gggegegee eegetetgeg etgtetetee g atg geg
                                                         Met Ala
tee gee tea ggg gee atg geg aag cae gag eag ate etg gte ete gat
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Ser Ala Ser Gly Ala Met Ala Lys His Glu Gln Ile Leu Val Leu Asp
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														cag Gln		477
														gta Val		525
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gct Ala	gtt Val	cca Pro	ctg Leu 150	aat Asn	gca Ala	tct Ser	aag Lys	caa Gln 155	gat Asp	gga Gly	cct Pro	atg Met	cca Pro 160	aaa Lys	cca Pro	717
cac His	agt Ser	gtt Val 165	tca Ser	ctt Leu	aat Asn	gat Asp	acc Thr 170	gaa Glu	aca Thr	agg Arg	aaa Lys	cta Leu 175	atg Met	gaa Glu	gag Glu	765
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														tcg Ser		861
														acc Thr 225		909
cct Pro	ctt Leu	cct Pro	tca Ser 230	ctt Leu	ctt Leu	gtt Val	gta Val	att Ile 235	gca Ala	gcc Ala	att Ile	ttc Phe	att Ile 240	gga Gly	ttc Phe	957

ttt cta ggg aaa ttc atc ttg tagagtgaag catgcagagt gctgtttctt Phe Leu Gly Lys Phe Ile Leu 245

1008

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Leu Arg Pro Glu His Phe Gln Glu Val Gly Tyr Ala Ala Pro Pro Ser 35 40 45

Pro Pro Leu Ser Arg Ser Leu Pro Met Asp His Pro Asp Ser Ser Gln 50 55 60

His Gly Pro Pro Phe Glu Gly Gln Ser Gln Val Gln Pro Pro Pro Ser 65 70 75 80

Gln Glu Ala Thr Pro Leu Gln Gln Glu Lys Leu Leu Pro Ala Gln Leu 85 90 95

Pro Ala Glu Lys Glu Val Gly Pro Pro Leu Pro Gl
n Glu Ala Val Pro 100 \$100\$

Leu Gln Lys Glu Leu Pro Ser Leu Gln His Pro Asn Glu Gln Lys Glu
115 120 125

Gly Thr Pro Ala Pro Phe Gly Asp Gln Ser His Pro Glu Pro Glu Ser

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Gly	His	Arg	Leu	Asp 165	Gly	Phe	Pro	Pro	Gly 170	Arg	Pro	Ser	Pro	Asp 175	Asr
Leu	Asn	Gln	Ile 180	Cys	Leu	Pro	Asn	Arg 185	Gln	His	Val	Val	Tyr 190	Gly	Pro
Trp	Asn	Leu 195	Pro	Gln	Ser	Ser	Tyr 200	Ser	His	Leu	Thr	Arg 205	Gln	Gly	Glu
Thr	Leu 210	Asn	Phe	Leu	Glu	Ile 215	Gly	Tyr	Ser	Arg	Cys 220	Cys	His	Cys	Arç
Ser 225	His	Thr	Asn		Leu 230	Glu	Cys	Ala	Lys	Leu 235	Val	Trp	Glu	Glu	Ala 240
Met	Ser	Arg	Phe	Cys 245	Glu	Ala	Glu	Phe	Ser 250	Val	Lys	Thr	Arg	Pro 255	His
Trp	Cys	Cys	Thr 260	Arg	Gln	Gly	Glu	Ala 265	Arg	Phe	Ser	Cys	Phe 270	Gln	Glu
Glu	Ala	Pro 275	Gln	Pro	His	Tyr	Gln 280	Leu	Arg	Ala	Суѕ	Pro 285	Ser	His	Glr
Pro	Asp 290	Ile	Ser	Ser	Gly	Leu 295	Glu	Leu	Pro	Phe	Pro 300	Pro	Gly	Val	Pro
Thr 305	Leu	Asp	Asn	Ile	Lys 310	Asn	Ile	Суѕ	His	Leu 315	Arg	Arg	Phe	Arg	Ser 320
Val	Pro	Arg	Asn	Leu 325	Pro	Ala	Thr	Asp	Pro 330	Leu	Gln	Arg	Glu	Leu 335	Leu
Ala	Leu	Ile	Gln 340	Leu	Glu	Arg	Glu	Phe 345	Gln	Arg	Cys	Cys	Arg 350	Gln	Gly
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Tyr	Cys 370	Asp	Arg	Glu	Tyr	Ala 375	Val	Lys	Thr	His	His 380	His	Leu	Суз	Cys
Arg 385	His	Pro	Pro	Ser	Pro 390	Thr	Arg	Asp	Glu	Cys 395	Phe	Ala	Arg	Arg	Ala 400
Pro	Tyr	Pro	Asn	Tyr 405	Asp	Arg	Asp	Ile	Leu 410	Thr	Ile	Asp	Ile	Gly 415	Arg
Val	Thr	Pro	Asn 420	Leu	Met	Gly	His	Leu 425	Cys	Gly	Asn	Gln	Arg 430	Val	Leu
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Cys	Cys 450	Asp	Leu	Pro	Phe	Pro 455	Glu	Gln	Ala	Cys	Cys 460	Ala	Glu	Glu	Glu.	
Lys 465	Leu	Thr	Phe	Ile	Asn 470	Asp	Leu	Cys	Gly	Pro 475	Arg	Arg	Asn	Ile	Trp 480	
Arg	Asp	Pro	Ala	Leu 485	Cys	Cys	Tyr	Leu	Ser 490	Pro	Gly	Asp	Glu	Gln 495	Val	
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	gca Ala															164
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	caa Gln						-								_	260
_	ctc Leu 55		_				_			_						308
	gga Gly															356

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ctt Leu	cct Pro	aac Asn	cgt Arg 185	cag Gln	cat His	gtg Val	gta Val	tat Tyr 190	ggt Gly	ccc Pro	tgg Trp	aac Asn	cta Leu 195	cca Pro	cag Gln	692
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						tgc Cys										1172
						gat Asp										1220
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ctactactca	cattcattta	atggtttctg	g taaacatttt	taagacagtt	gggatgtcac 2106

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Arg Val Gln Gln Asn Val Pro Ser Gly Thr Asp Thr Gly Asp Pro Gln 35 40 45

Ser Lys Pro Leu Gly Asp Trp Ala Ala Gly Thr Met Asp Pro Glu Ser 50 55 60

Ser Ile Phe Ile Glu Asp Ala Ile Lys Tyr Phe Lys Glu Lys Val Ser 65 70 75 80

- Thr Gln Asn Leu Leu Leu Leu Thr Asp Asn Glu Ala Trp Asn Gly 85 90 95
- Phe Val Ala Ala Ala Glu Leu Pro Arg Asn Glu Ala Asp Glu Leu Arg 100 105 110
- Lys Ala Leu Asp Asn Leu Ala Arg Gln Met Ile Met Lys Asp Lys Asn 115 120 125
- Trp His Asp Lys Gly Gln Gln Tyr Arg Asn Trp Phe Leu Lys Glu Phe 130 135 140
- Pro Arg Leu Lys Ser Lys Leu Glu Asp Asn Ile Arg Arg Leu Arg Ala 145 150 150 160
- Leu Ala Asp Gly Val Gln Lys Val His Lys Gly Thr Thr Ile Ala Asn 165 170 175
- Val Val Ser Gly Ser Leu Ser Ile Ser Ser Gly Ile Leu Thr Leu Val 180 185 190
- Gly Met Gly Leu Ala Pro Phe Thr Glu Gly Gly Ser Leu Val Leu Leu 195 200 205
- Glu Pro Gly Met Glu Leu Gly Ile Thr Ala Ala Leu Thr Gly Ile Thr 210 215 220
- Ser Ser Thr Ile Asp Tyr Gly Lys Lys Trp Trp Thr Gln Ala Gln Ala 225 230 235 240
- His Asp Leu Val Ile Lys Ser Leu Asp Lys Leu Lys Glu Val Lys Glu 245 250 255
- Phe Leu Gly Glu Asn Ile Ser Asn Phe Leu Ser Leu Ala Gly Asn Thr 260 265 270
- Tyr Gln Leu Thr Arg Gly Ile Gly Lys Asp Ile Arg Ala Leu Arg Arg 275 280 285
- Ala Arg Ala Asn Leu Gln Ser Val Pro His Ala Ser Ala Ser Arg Pro 290 295 300
- Arg Val Thr Glu Pro Ile Ser Ala Glu Ser Gly Glu Gln Val Glu Arg 305 310 315 320
- Val Asn Glu Pro Ser Ile Leu Glu Met Ser Arg Gly Val Lys Leu Thr 325 330 335
- Asp Val Ala Pro Val Ser Phe Phe Leu Val Leu Asp Val Val Tyr Leu 340 345 350
- Val Tyr Glu Ser Lys His Leu His Glu Gly Ala Lys Ser Glu Thr Ala 355 360 365
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Leu Asn Asn Asn Tyr Lys Ile Leu Gln Ala Asp Gln Glu Leu

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ctg Leu	acc Thr 190	ctc Leu	gtc Val	ggc Gly	atg Met	ggt Gly 195	ctg Leu	gca Ala	ccc Pro	ttc Phe	aca Thr 200	gag Glu	gga Gly	ggc	agc Ser	687
ctt Leu 205	gta Val	ctc Leu	ttg Leu	gaa Glu	cct Pro 210	ggg	atg Met	gag Glu	ttg Leu	gga Gly 215	atc Ile	aca Thr	gca Ala	gct Ala	ttg Leu 220	735
acc Thr	ggg Gly	att Ile	acc Thr	agc Ser 225	agt Ser	acc Thr	ata Ile	gac Asp	tac Tyr 230	gga Gly	aag Lys	aag Lys	tgg Trp	tgg Trp 235	aca Thr	783
caa Gln	gcc Ala	caa Gln	gcc Ala 240	cac His	gac Asp	ctg Leu	gtc Val	atc Ile 245	aaa Lys	agc Ser	ctt Leu	gac Asp	aaa Lys 250	ttg Leu	aag Lys	831
gag Glu	gtg Val	aag Lys 255	gag Glu	ttt Phe	ttg Leu	ggt Gly	gag Glu 260	aac Asn	ata Ile	tcc Ser	aac Asn	ttt Phe 265	ctt Leu	tcc Ser	tta Leu	879
gct Ala	ggc Gly 270	aat Asn	act Thr	tac Tyr	caa Gln	ctc Leu 275	aca Thr	cga Arg	ggc	att Ile	ggg Gly 280	aag Lys	gac Asp	atc Ile	cgt Arg	927
gcc Ala 285	ctc Leu	aga Arg	cga Arg	gcc Ala	aga Arg 290	gcc Ala	aat Asn	ctt Leu	cag Gln	tca Ser 295	Val	ccg Pro	cat His	gcc Ala	tca Ser 300	975
gcc Ala	tca Ser	cgc Arg	ccc Pro	cgģ Arg 305	Val	act Thr	gag Glu	cca Pro	ato Ile 310	Ser	gct Ala	gaa Glu	agc Ser	ggt Gly 315	Glu	1023
cag Gln	gtg Val	gag Glu	aga Arg 320	Val	aat Asn	gaa Glu	ccc Pro	ago Ser 325	Ile	ctg Leu	gaa Glu	atg Met	ago Ser 330	Arg	gga Gly	1071
gtc Val	aag Lys	cto Leu 335	Thr	gat Asp	gtg Val	gcc Ala	e cct Pro 340	Val	ago Ser	ttc Phe	: ttt : Phe	ctt Leu 345	. Val	r ctg Leu	gat Asp	1119
gta Val	gtc Val	Tyr	ctc Leu	gtg Val	tac Tyr	gaa Glu 355	ı Ser	aag Lys	cac His	tta Leu	cat His 360	Glu	r GJZ L ggg	g gca ⁄ Ala	aag Lys	1167
tca Ser 365	Glu	g aca Thr	a gct Ala	gaç Glu	gag Glu 370	Let	g aag Lys	g aag S Lys	g gto Val	g gct Ala 375	a Glr	g gag n Glu	g cto Lei	g gag ı Glu	gag Glu 380	1215
aaq Lys	g cta Leu	aad Asr	att n Ile	cto Lev	c aac ı Asr	aat Asr	aat Asr	tat Tyr	aaq Lys	g att	cto e Lei	g caq ı Glr	g gcg n Ala	g gac a Asp	caa Gln	1263

385 390 395

gaa ctg tgaccacagg gcagggcagc caccaggaga gatatgcctg gcaggggcca 1319 Glu Leu ggacaaaatg caaacttttt tttttctga gacagagtct tgctctgtcg ccaagttgca 1379 gtgagccgag atatcgccac tgcactccag cctgggtgac agagcgagac tccatctcaa 1439 aaaaaaaaaa aaaaagaata tattgacgga agaatagaga ggaggcttga aggaaccagc 1499 aatgagaagg ccaggaaaag aaagagctga aaatggagaa agcccaagag ttagaacagt 1559 tggatacagg agaagaaaca geggeteeae tacagâceea geeecaggtt caatgteete 1619 cgaagaatga agtettteee tggtgatggt eccetgeeet gtettteeag cateeactet 1679 cccttgtcct cctgggggca tatctcagtc aggcagcggc ttcctgatga tggtcgttgg 1739 ggtggttgtc atgtgatggg tcccctccag gttactaaag ggtgcatgtc ccctgcttga 1799 acactgaagg gcaggtggtg agccatggcc atggtcccca gctgaggagc aggtgtccct 1859 gagaacccaa acttcccaga gagtatgtga gaaccaacca atgaaaacag tcccatcgct 1919 cttacccggt aagtaaacag tcagaaaatt agcatgaaag cagtttagca ttgggaggaa 1979 getcagatet etagagetgt ettgteeceg eccaggattg acetgtgtaa gteecaataa 2039 2054 actcacctac tcatc

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<212> PRT

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<400> 119

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Arg Val Gln Gln Asn Val Pro Ser Gly Thr Asp Thr Gly Asp Pro Gln
35 40 45

Ser Lys Pro Leu Gly Asp Trp Ala Ala Gly Thr Met Asp Pro Glu Ser 50 55 60

Ser Ile Phe Ile Glu Asp Ala Ile Lys Tyr Phe Lys Glu Lys Val Ser 65 70 75 80

Thr Gln Asn Leu Leu Leu Leu Thr Asp Asn Glu Ala Trp Asn Gly 85 90 95

Phe Val Ala Ala Ala Glu Leu Pro Arg Asn Glu Ala Asp Glu Leu Arg

			100					105					110		
Lys	Ala	Leu 115	Asp	Asn	Leu	Ala	Arg 120	Gln	Met	Ile	Met	Lys 125	Asp	Lys	Asn
Trp	His 130	Asp	Lys	Gly	Gln	Gln 135	Tyr	Arg	Asn	Trp	Phe 140	Leu	Lys	Glu	Phe
Pro 145	Arg	Leu	Lys	Ser	Lys 150	Leu	Glu	Asp	Asn	Ile 155	Arg	Arg	Leu	Arg	Ala 160
Leu	Ala	Asp	Gly	Val 165	Gln	Lys	Val	His	Lys 170	Gly	Thr	Thr	Ile	Ala 175	Asn
Val	Val	Ser	Gly 180	Ser	Leu	Ser	Ile	Ser 185	Ser	Gly	Ile	Leu	Thr 190	Leu	Val
Gly	Met	Gly 195	Leu	Ala	Pro	Phe	Thr 200	Glu	Gly	Gly	Ser	Leu 205	Val	Leu	Leu
Glu	Pro 210	Gly	Met	Glu	Leu	Gly 215	Ile	Thr	Ala	Ala	Leu 220	Thr	Gly	Ile	Thr
Ser 225	Ser	Thr	Ile	Asp	Tyr 230	Gly	Lys	Lys	Trp	Trp 235	Thr	Gln	Ala	Gln	Ala 240
His	Asp	Leu	Val	Ile 245		Ser	Leu	Asp	Lys 250	Leu	Lys	Glu	Val	Lys 255	Glu
Phe	Leu	Gly	Glu 260		Ile	Ser	Asn	Phe 265	Leu	Ser	Leu	Ala	Gly 270	Asn	Thr
Tyr	Gln	Leu 275		Arg	Gly	Ile	Gly 280		Asp	Ile	Arg	Ala 285		Arg	Arg
Ala	Arg 290		Asn	Leu	Gln	Ser 295		Pro	His	Ala	Ser 300		Ser	Arg	Pro
Arg 305		Thr	Glu	Pro	310		Ala	Glu	Ser	Gly 315		Gln	Val	Glu	Arg 320
Val	Asn	Glu	Pro	Ser 325		Leu	Glu	Met	Ser 330		Gly	Val	Lys	Leu 335	
Asp	Val	Ala	Pro 340		Ser	Phe	Phe	Leu 345	Val	Leu	Asp	Val	Val 350		Lev
Val	Tyr	Glu 355		Lys	His	Leu	His 360		Gly	Ala	Lys	Ser 365		Thr	Ala
Glu	Glu 370		ı Lys	Lys	s Val	Ala 375		Glu	l Leu	Glu	380		. Leu	. Asn	ıIl∈
Leu 385		ı Asr	n Asr	туг	Lys 390		e Leu	ı Glr	n Ala	Asp 395		Glu	ı Leu	L .	

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tgc atc tgg atg agt gca ctt ttc ctt ggt gtg aga gtg agg gca gag
Cys Ile Trp Met Ser Ala Leu Phe Leu Gly Val Arg Val Arg Ala Glu
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gaa gct gga gcg agg gtg caa caa aac gtt cca agt ggg aca gat act
Glu Ala Gly Ala Arg Val Gln Gln Asn Val Pro Ser Gly Thr Asp Thr
                          35
     30
gga gat cct caa agt aag ccc ctc ggt gac tgg gct gct ggc acc atg
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Gly Asp Pro Gln Ser Lys Pro Leu Gly Asp Trp Ala Ala Gly Thr Met
                                          5.5
 45
                                                                    303
gac cca gag age agt atc ttt att gag gat gcc att aag tat ttc aag
Asp Pro Glu Ser Ser Ile Phe Ile Glu Asp Ala Ile Lys Tyr Phe Lys
                                      7.0
                  65
gaa aaa gtg agc aca cag aat ctg cta ctc ctg ctg act gat aat gag
                                                                    351
Glu Lys Val Ser Thr Gln Asn Leu Leu Leu Leu Leu Thr Asp Asn Glu
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gcc tgg aac gga ttc gtg gct gct gct gaa ctg ccc agg aat gag gca
                                                                    399
Ala Trp Asn Gly Phe Val Ala Ala Ala Glu Leu Pro Arg Asn Glu Ala
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                                                                    447
gat gag etc egt aaa get etg gae aac ett gea aga eaa atg ate atg
Asp Glu Leu Arg Lys Ala Leu Asp Asn Leu Ala Arg Gln Met Ile Met
    110
                         115
                                                                    495
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Lys Asp Lys Asn Trp His Asp Lys Gly Gln Gln Tyr Arg Asn Trp Phe
                                                              140
                     130
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 ctg aaa gag ttt cct cgg ttg aaa agt aag ctt gag gat aac ata aga
                                                                    543
Leu Lys Glu Phe Pro Arg Leu Lys Ser Lys Leu Glu Asp Asn Ile Arg
                                                          155
                 145
 agg ctc cgt gcc ctt gca gat ggg gtt cag aag gtc cac aaa ggc acc
 Arg Leu Arg Ala Leu Ala Asp Gly Val Gln Lys Val His Lys Gly Thr
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             160
 acc atc gcc aat gtg gtg tct ggc tct ctc agc att tcc tct ggc atc
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ctt Leu 205	gta Val	ctc Leu	ttg Leu	gaa Glu	cct Pro 210	ggg Gly	atg Met	gag Glu	ttg Leu	gga Gly 215	atc Ile	aca Thr	gca Ala	gct Ala	ttg Leu 220	735
acc Thr	Gly ggg	att Ile	acc Thr	agc Ser 225	agt Ser	acc Thr	ata Ile	gac Asp	tac Tyr 230	gga Gly	aag Lys	aag Lys	tgg Trp	tgg Trp 235	aca Thr	783
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gcc Ala	tca Ser	cgc Arg	ccc Pro	cgg Arg 305	Val	act Thr	gag Glu	cca Pro	atc Ile 310	Ser	gct Ala	gaa Glu	agc Ser	ggt Gly 315	gaa Glu	1023
caç Glr	gtg Val	.gag Glu	aga Arg 320	y Val	aat Asn	gaa Glu	ccc Pro	agc Ser 325	Ile	ctg Leu	gaa Glu	atg Met	ago Ser 330	Arg	gga Gly	1071
gto Val	aag Lys	cto Lev 335	ı Thr	gat Asp	gtg Val	gcc Ala	cct Pro	Val	ago Ser	ttc Phe	ttt Phe	ctt Leu 345	ı Val	ctç Lev	gat Asp	1119
gta Val	a gtc L Val 350	. Туз	c cto Lei	gtç ı Val	tac Tyr	gaa Glu 355	Ser	aag Lys	g cac His	tta Lei	a cat ı His 360	s Glu	ı Gl? I ggç	g gca 7 Ala	a aag a Lys	1167
tca Sei 36!	c Glu	g aca	a gct r Ala	z gaç a Glı	g gag ı Glu 370	ı Lev	g aag Lys	g aag s Lys	g gto s Val	g gct L Ala 375	a GIr	g gaç n Glu	g cto i Lei	g gaq ı Glı	g gag 1 Glu 380	1215
aa Ly:	g cta s Lei	a aad 1 Asi	c att	t cto e Leo 385	ג Asr	aat n Asr	aat n Asr	tat n Tym	aaq Lys 390	s Ile	t cto	g caq ı Glı	g gco n Ala	g gad a Ası 39!	c caa p Gln	1263
	a cto u Leo		acca	cagg	gca	gggca	agc (cacca	agga	ga ga	atato	geet	ggc	aggg	gcca	1319

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<211> 108

<212> PRT

<213> Homo sapiens

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Asp Gly Lys Lys Phe Asp Ser Ser Arg Asp Arg Asn Lys Pro Phe Lys

Phe Met Leu Gly Lys Gln Glu Val Ile Arg Gly Trp Glu Glu Gly Val 50 60

Ala Gln Met Ser Val Gly Gln Arg Ala Lys Leu Thr Ile Ser Pro Asp
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Tyr Ala Tyr Gly Ala Thr Gly His Pro Gly Ile Ile Pro Pro His Ala 85 90 95

Thr Leu Val Phe Asp Val Glu Leu Leu Lys Leu Glu 100 105

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tiggeggettt tete 1546

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<211> 679

<212> PRT

<213> Homo sapiens

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Phe Val Leu Ala Phe Ser Val Gly Ala Asn Asp Val Ala Asn Ser Phe 35 40 45

Gly Thr Ala Val Gly Ser Gly Val Val Thr Leu Lys Gln Ala Cys Ile 50 55 60

Leu Ala Ser Ile Phe Glu Thr Val Gly Ser Val Leu Leu Gly Ala Lys
65 70 75 80

Val Ser Glu Thr Ile Arg Lys Gly Leu Ile Asp Val Glu Met Tyr Asn 85 90 95

Ser Thr Gln Gly Leu Leu Met Ala Gly Ser Val Ser Ala Met Phe Gly 100 105 110

Ser Ala Val Trp Gln Leu Val Ala Ser Phe Leu Lys Leu Pro Ile Ser 115 120 125

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caa Gl:	a geo n Ala 365	. Val	ago Ser	aac Asn	caa Gln	ata Ile 370	Asn	tcc Ser	agt Ser	ggc Gly	cac His	Tyr	cag Gln	tat Tyr	cac His	1217
acc Thi	. Val	g cat His	aag Lys	g gat s Asp	tcc Ser 385	· Gly	ctg Leu	tac Tyr	aaa Lys	gag Glu 390	ı Let	cto Leu	cat His	aaa Lys	tta Leu 395	1265
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Ile Met Ala Ile Phe Arg Tyr Val Ile Leu Arg Met 670 675

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Ile Glu Arg Asn Lys Glu Ser Pro Val Phe Ala Pro Val Tyr Phe Pro 65 70 75 80

Glu Glu Leu His Arg Lys Ala Ala Leu Glu Gln Asp Leu Ala Phe Trp 85 90 95

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Gln 145	Val	Leu	Lys	Lys	Ile 150	Ala	Gln	Lys	Ala	Leu 155	Asp	Leu	Pro	Ser	Ser 160	
Gly	Glu	Gly	Leu	Ala 165	Phe	Phe	Thr	Phe	Pro 170	Asn	Ile	Ala	Ser	Ala 175	Thr	
Lys	Phe	Lys	Gln 180	Leu	Tyr	Arg	Ser	Arg 185	Met	Asn	Ser	Leu	Glu 190	Met	Thr	
Pro	Ala	Val 195	Arg	Gln	Arg	Val	Ile 200	Glu	Glu	Ala	Lys	Thr 205	Ala	Phe	Leu .	
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Thr 225	Lys	Asp	Gln	Ser	Pro 230	Ser	Arg	Ala	Pro	Gly 235		Arg	Gln	Arg	Ala 240	
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Pro	Pro	Leu	Asn 260	Thr	Arg	Ser	Gln	Ala 265	Pro	Leu	Leu	Arg	Trp 270		Leu	
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															ctt Leu	881

255 260 265

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Arg Phe Asn Ala His Gly Asp Ala Asn Thr Ile Val Cys Asn Ser Lys 50 55 60

Asp Gly Gly Ala Trp Gly Thr Glu Gln Arg Glu Ala Val Phe Pro Phe 65 70 75 80

Gln Pro Gly Ser Val Ala Glu Val Cys Ile Thr Phe Asp Gln Ala Asn 85 90 95

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Gly Lys Asp Ser Asn Asn Leu Cys Leu His Phe Asn Pro Arg Phe Asn
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 Glu Ala Ile Asn Tyr Met Ala Ala Asp Gly Asp Phe Lys Ile Lys Cys
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Thr Leu Val Leu Gln Cys Val Ala Val Gly Ser Pro Ile Pro His Tyr
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Gln Trp Phe Lys Asn Glu Leu Pro Leu Thr His Glu Thr Lys Lys Leu 100 105 110

Tyr Met Val Pro Tyr Val Asp Leu Glu His Gln Gly Thr Tyr Trp Cys 115 120 125

His Val Tyr Asn Asp Arg Asp Ser Gln Asp Ser Lys Lys Val Glu Ile 130 135 140

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tca Sei	a gga c Gly	tto Lev 695	ı Glu	a gat ı Asp	act Thr	gta Val	a gaç L Glu 700	ı Asp	c aaq o Lys	g caç s Glr	g gaa n Glu	a gto u Val 705	Asr	gtt Val	r Glà	2288
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Cys Glu Val Ser 50	His Arg Are		Gly Trp Asn Glu Phe Asp	
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Asn Pro Leu Ile Met Leu Leu Leu Ala Ser Ala Val Ile Ser Val Leu

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Glu	Glu 130	Leu	Ser	Lys	Leu	Val 135	Pro	Pro	Glu	Суѕ	His 140	Cys	Val	Arg	Glu
Gly 145	Lys	Leu	Glu	His	Thr 150	Leu	Ala	Arg	Asp	Leu 155	Val	Pro	Gly	Asp	Thr 160
Val	Cys	Leu	Ser	Val 165	Gly	Asp	Arg	Val	Pro 170	Ala	Asp	Leu	Arg	Leu 175	Phe
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Phe	Gly	Glu	Val	Phe 245	Lys	Met	Met	Gln	Ala 250		. Glu	Ala	Pro	Lys 255	Thr
Pro	Leu	Gln	Lys 260		Met	Asp	Leu	Leu 265		, Lys	Gln	Leu	Ser 270	Phe	Tyr
Ser	Phe	Gly 275		lle	Gly	Ile	280		Leu	ı Val	Gly	7 Trp 285	Leu	Leu	Gly
Lys	Asp 290		Leu	Glu	Met	Phe 295		· Ile	Ser	Val	Ser 300	Leu	ı Ala	. Val	Ala
Ala 305		Pro	Glu	ı Gly	Leu 310		ıle	· Val	. Val	Thr 315	Val	. Thr	Lev	Ala	Let 320
Gly	v Val	. Met	. Arg	Met 325	Val	Lys	. Lys	: Arg	330		e Val	Lys	Lys	335	Pro
Ile	val	. Glu	Thr 340		ı Gly	cys	суя	345		l Ile	e Cys	s Ser	350	Lys)	Thi
Gly	7 Thr	Leu 355		c Lys	s Asn	Glu	1 Met 360		val	l Thi	c His	365	e Phe	e Thr	Sei
Asp	Gl _y 370		ı His	s Ala	a Glu	val 375		Gly	y Val	l Gl	у Туя 380		n Glr	n Phe	e Gly

Glu Val Ile Val Asp Gly Asp Val Val His Gly Phe Tyr Asn Pro Ala

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Ser	· Val	Ser 595		Glu	ı Glu	Ile	Asp 600	Ala	Met	Asp	Val	Gln 605	Gln	Leu	Ser
Glr	11e		Pro	Lys	s Val	Ala 615		Phe	Туг	Arg	Ala 620	Ser	Pro	Arg	His
Lys 625		Lys	s Ile	e Ile	E Lys 630		Leu	Gln	Lys	635	Gly	Ser	· Val	. Val	Ala 640
Met	Th:	c Gly	y Asp	Gl ₃ 645		Asr	a Asp	Ala	Val 650	L Ala	Leu	Lys	ala Ala	Ala 655	a Asp
Ile	e Gly	y Val	L Ala		c Gly	/ Glr	n Thr	G13		r Asp	o Val	. Cys	670	s Glu)	ı Ala
Ala	a Ası	p M e 67		e Let	ı Val	L Asp	Asp 680) Phe	e Glr	n Thr	11e 685	e Met	sei	r Ala
Il	e Gl	u Gl	u Gl	v Lv:	s Gly	y Ile	э Туз	. Ası	n Ası	n Ile	e Lys	s Asr	n Phe	e Val	l Arg

695 700 690 Phe Gln Leu Ser Thr Ser Ile Ala Ala Leu Thr Leu Ile Ser Leu Ala 710 Thr Leu Met Asn Phe Pro Asn Pro Leu Asn Ala Met Gln Ile Leu Trp 725 Ile Asn Ile Ile Met Asp Gly Pro Pro Ala Gln Ser Leu Gly Val Glu Pro Val Asp Lys Asp Val Ile Arg Lys Pro Pro Arg Asn Trp Lys Asp 760 Ser Ile Leu Thr Lys Asn Leu Ile Leu Lys Ile Leu Val Ser Ser Ile 775 Ile Ile Val Cys Gly Thr Leu Phe Val Phe Trp Arg Glu Leu Arg Asp Asn Val Ile Thr Pro Arg Asp Thr Thr Met Thr Phe Thr Cys Phe Val 810 Phe Phe Asp Met Phe Asn Ala Leu Ser Ser Arg Ser Gln Thr Lys Ser 825 Val Phe Glu Ile Gly Leu Cys Ser Asn Arg Met Phe Cys Tyr Ala Val 840 Leu Gly Ser Ile Met Gly Gln Leu Leu Val Ile Tyr Phe Pro Pro Leu 850 Gln Lys Val Phe Gln Thr Glu Ser Leu Ser Ile Leu Asp Leu Leu Phe 875 Leu Leu Gly Leu Thr Ser Ser Val Cys Ile Val Ala Glu Ile Ile Lys Lys Val Glu Arg Ser Arg Glu Lys Ile Gln Lys His Val Ser Ser Thr 905 Ser Ser Ser Phe Leu Glu Val 915 <210> 134 <211> 3612 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (427)..(3183) <400> 134 acggcctcgc ggagccggcc cggcggaccg tgacgggtcc cctcacctcc tcttctccc 60

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Thr Leu Phe Val Asn Gly Gln Pro Arg Pro Leu Glu Ser Ser Gln Val
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			cag cca aga o Gln Pro Arg 1 55										
			ctg ata gaa d Leu Ile Glu I										
	Leu Leu Asp		cca cct gga (Pro Pro Gly (
			gtg gat ggt a Val Asp Gly 2 105										
			tct act cag Ser Thr Gln	Val Met Ala									
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			gat gat cag Asp Asp Gln										
	Pro Ala Glu		gga aca ccc Gly Thr Pro										
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gcc Ala	tct Ser	caa Gln	cct Pro 305	gga Gly	atg Met	gct Ala	cca Pro	agc Ser 310	caa Gln	cct Pro	ggg	gcc Ala	tat Tyr 315	caa Gln	cca Pro	961
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agt Ser	ggg Gly 335	Pro	aat Asn	cct Pro	tat Tyr	gcg Ala 340	Arg	aac Asn	cgt Arg	cct Pro	ccc Pro 345	Phe	ggt Gly	cag Gln	22	·1057
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Leu Ser Asn Asp Glu Val Thr Ile Lys Tyr Lys Asp Glu Asp Gly Asp 50 60

Leu Ile Thr Ile Phe Asp Ser Ser Asp Leu Ser Phe Ala Ile Gln Cys
65 70 75 80

Ser Arg Ile Leu Lys Leu Thr Leu Phe Val Asn Gly Gln Pro Arg Pro 85 90 95

Leu Glu Ser Ser Gln Val Lys Tyr Leu Arg Arg Glu Leu Ile Glu Leu 100 105 110

Arg Asn Lys Val Asn Arg Leu Leu Asp Ser Leu Glu Pro Pro Gly Glu 115 120 125

Pro Gly Pro Ser Thr Asn Ile Pro Glu Asn Asp Thr Val Asp Gly Arg 130 135

Glu Glu Lys Ser Ala Ser Asp Ser Ser Gly Lys Gln Ser Thr Gln Val 145 150 155 160

Met Ala Ala Ser Met Ser Ala Phe Asp Pro Leu Lys Asn Gln Asp Glu 165 170 175

Ile Asn Lys Asn Val Met Ser Ala Phe Gly Leu Thr Asp Asp Gln Val 180 185 190

Ser Gly Pro Pro Ser Ala Pro Ala Glu Asp Arg Ser Gly Thr Pro Asp 195 200 205

Ser Ile Ala Ser Ser Ser Ser Ala Ala His Pro Pro Gly Val Gln Pro 210 215 220

Gln Gln Pro Pro Tyr Thr Gly Ala Gln Thr Gln Ala Gly Gln Met Tyr 225 230 235

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Pro Pro Gln Gln Pro Gln Gln Tyr Gly Ile Gln Tyr Ser Ala Ser Tyr 260 265 270

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Tyr	Gln	Pro 355	Arg	Pro	Gly	Phe	Thr 360	Ser	Leu	Pro	Gly	Ser 365	Thr	Met	Thr	
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				act Thr												147
				ctt Leu												195
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Val	Asn	Gly 35		. Val	Leu	Ser	Ser 40		Gly	Thr	Arg	Phe 45		Val	Asn	
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His Arg Val Asp Thr Ile Ser Val Asn Gly Ser Val Gln Leu Ser Tyr 130 135 140

Thr Gln Thr Val Ile His Thr Val Gln Ser Ala Pro Gly Gln Met Phe 165 170 175

Ser Thr Pro Ala Ile Pro Pro Met Met Tyr Pro His Pro Ala Tyr Pro 180 185 190

Met Pro Phe Ile Thr Thr Ile Leu Gly Gly Leu Tyr Pro Ser Lys Ser 195 200 205

Ile Leu Leu Ser Gly Thr Val Leu Pro Ser Ala Gln Arg Phe His Ile 210 215 220

Asn Leu Cys Ser Gly Asn His Ile Ala Phe His Leu Asn Pro Arg Phe 225 230 235 240

Asp Glu Asn Ala Val Val Arg Asn Thr Gln Ile Asp Asn Ser Trp Gly 245 250 255

Ser Glu Glu Arg Ser Leu Pro Arg Lys Met Pro Phe Val Arg Gly Gln 260 265 270

Ser Phe Ser Val Trp Ile Leu Cys Glu Ala His Cys Leu Lys Val Ala 275 280 285

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	_	_	aat Asn	-					_							524
_	_		agc Ser	_			_	-						_		572
_	_		ctg Leu 120	_	_	_				_		_	-			620
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<213> Homo sapiens

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- Ile Arg Glu Asn Ala Glu Val Ser Met Asp Val Ser Leu Ala Tyr Arg 50 55 60
- Asp Asp Ala Phe Ala Glu Trp Thr Glu Met Ala His Glu Arg Val Pro 65 70 75 80
- Arg Lys Leu Lys Cys Thr Phe Thr Ser Pro Lys Thr Pro Glu His Glu 85 90 95
- Gly Arg Tyr Tyr Glu Cys Asp Val Leu Pro Phe Met Glu Ile Gly Ser 100 105 110
- Val Ala His Lys Phe Tyr Leu Leu Asn Ile Arg Leu Pro Val Asn Glu 115 120 125
- Lys Lys Ile Asn Val Gly Ile Gly Glu Ile Lys Asp Ile Arg Leu 130 135 140
- Val Gly Ile His Gln Asn Gly Gly Phe Thr Lys Val Trp Phe Ala Met 145 \$150\$ 150 \$155\$
- Lys Thr Phe Leu Thr Pro Ser Ile Phe Ile Ile Met Val Trp Tyr Trp 165 170 175
- Arg Arg Ile Thr Met Met Ser Arg Pro Pro Val Leu Leu Glu Lys Val 180 185 190
- Ile Phe Ala Leu Gly Ile Ser Met Thr Phe Ile Asn Ile Pro Val Glu 195 200 205
- Trp Phe Ser Ile Gly Phe Asp Trp Thr Trp Met Leu Leu Phe Gly Asp 210 215 220
- Ile Arg Gln Gly Ile Phe Tyr Ala Met Leu Leu Ser Phe Trp Ile Ile 225 230 235
- Phe Cys Gly Glu His Met Met Asp Gln His Glu Arg Asn His Ile Ala 245 250 255
- Gly Tyr Trp Lys Gln Val Gly Pro Ile Ala Val Gly Ser Phe Cys Leu 260 265 270
- Phe Ile Phe Asp Met Cys Glu Arg Gly Val Gln Leu Thr Asn Pro Phe 275 280 285
- Tyr Ser Ile Trp Thr Thr Asp Ile Gly Thr Glu Leu Ala Met Ala Phe 290 295 300

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Trp Gly Gly Val Thr Val Gln Val Asn Ser Ala Phe Phe Thr Gly Ile 385 390 395 400

Tyr Gly Met Trp Asn Leu Tyr Val Phe Ala Leu Met Phe Leu Tyr Ala 405 410 415

Pro Ser His Lys Asn Tyr Gly Glu Asp Gln Ser Asn Gly Met Gln Leu 420 425 430

Pro Cys Lys Ser Arg Glu Asp Cys Ala Leu Phe Val Ser Glu Leu Tyr 435 440 445

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		-		_	-	-	-		-	gag Glu 70			_	_	~	603
	_	_	_						_	acc Thr					_	651
									-	tgt Cys	-	-				699
	_					_				tac Tyr						747
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										ccc Pro						891
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				_						ttt Phe	-					1035
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Thr Phe Ile Asn Ile Pro Val Glu Trp Phe Ser Ile Gly Phe Asp Trp 195 200 205

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210 215 220
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Gly Thr Glu Leu Ala Met Ala Phe Ile Ile Val Ala Gly Ile Cys Leu 290 295 300

Cys Leu Tyr Phe Leu Phe Leu Cys Phe Met Val Phe Gln Val Phe Arg 305 310 315 320

Asn Ile Ser Gly Lys Gln Ser Ser Leu Pro Ala Met Ser Lys Val Arg 325 330 335

Arg Leu His Tyr Glu Gly Leu Ile Phe Arg Phe Lys Phe Leu Met Leu 340 345 350

Ile Thr Leu Ala Cys Ala Ala Met Thr Val Ile Phe Phe Ile Val Ser 355 360 365

Gln Val Thr Glu Gly His Trp Lys Trp Gly Gly Ile Thr Val Gln Val 370 375 380

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ttc co Phe Pr 45															555
gct ga Ala Gl															603
tgc ac Cys Th															651
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acg co						_				_					891

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			atg Met													987
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			gcg Ala													1083
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	_		ccc Pro		_	_				_						1179
-	_		aga Arg		-											1227
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			ctc Leu													1323
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Ser Val		ly Ala 00	Leu	Leu	Leu	Tyr 105	Met	Ala	Phe	Leu	Met 110	Leu	Val	
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Asp Arg His Lys Met Leu Ser
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				g ctt ctc cat s Leu Leu His	_
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			-		cat His	_		-	-	_	_		_			1430
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gcc cag t Ala Gln S						_				_	_	2342
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_	-	-		_	-	_	_					gtc Val				834
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		_		_		_				_	ccc Pro 530				_	1698
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	_	_			-			_	_		ttg Leu			_	-	1842
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Ser Val Glu Asp Thr Thr Ser Pro Asn Thr Lys Pro Cys Pro Pro Thr Pro Thr Thr Pro Glu Thr Ser Pro Pro Pro Pro Pro Pro Pro Ser Ser Thr Pro Cys Ser Ala His Leu Thr Pro Ser Ser Leu Phe Pro Ser 375 Ser Leu Glu Ser Ser Glu Gln Lys Phe Tyr Asn Phe Val Ile Leu 390 395 His Ala Arg Ala Asp Glu His Ile Ala Leu Arg Val Arg Glu Lys Leu 410 Glu Ala Leu Gly Val Pro Asp Gly Ala Thr Phe Cys Glu Asp Phe Gln 425 Val Pro Gly Arg Gly Glu Leu Ser Cys Leu Gln Asp Ala Ile Asp His Ser Ala Phe Ile Ile Leu Leu Thr Ser Asn Phe Asp Cys Arg Leu 455 Ser Leu His Gln Val Asn Gln Ala Met Met Ser Asn Leu Thr Arg Gln Gly Ser Pro Asp Cys Val Ile Pro Phe Leu Pro Leu Glu Ser Ser Pro Ala Gln Leu Ser Ser Asp Thr Ala Ser Leu Leu Ser Gly Leu Val Arg Leu Asp Glu His Ser Gln Ile Phe Ala Arg Lys Val Ala Asn Thr Phe 515 520 Lys Pro His Arg Leu Gln Ala Arg Lys Ala Met Trp Arg Lys Glu Gln 535 Asp Thr Arg Ala Leu Arg Glu Gln Ser Gln His Leu Asp Gly Glu Arg Met Gln Ala Ala Leu Asn Ala Ala Tyr Ser Ala Tyr Leu Gln Ser 570 Tyr Leu Ser Tyr Gln Ala Gln Met Glu Gln Leu Gln Val Ala Phe Gly Ser His Met Ser Phe Gly Thr Gly Ala Pro Tyr Gly Ala Arg Met Pro 600 Phe Gly Gly Gln Val Pro Leu Gly Ala Pro Pro Pro Phe Pro Thr Trp Pro Gly Cys Pro Gln Pro Pro Pro Leu His Ala Trp Gln Ala Gly Thr 630 635

Phe Pro Gln Ser Pro Ala Phe Pro Thr Ala Ser Pro Ala Pro Pro Gln 665 Ser Pro Gly Leu Gln Pro Leu Ile Ile His His Ala Gln Met Val Gln 680 Leu Gly Leu Asn Asn His Met Trp Asn Gln Arg Gly Ser Gln Ala Pro 695 700 Glu Asp Lys Thr Gln Glu Ala Glu 710 <210> 155 <211> 3456 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (281)..(3016) <400> 155 agcattcaag gagctcccca ggagaaagag caagttctga ggagccctct gagcccggaa 60 cgtgtccacc cggtcatgcc cgccgcgcac cagccccgca gtggacttgg aggaggagga 120 ggaggagage tetgtggatg geaaagggga eeggaagage acaggeetga aactetecaa 180 qaaqaaaqca aqqaqqaqac acacqqatqa cccaaqcaaq qaatqcttca ctctqaaatt 240 tgacctgaat gtggacattg agacagagat cgtcccagcc atg aag aag tca 295 Met Lys Lys Lys Ser 343 ctg ggg gag gtg ctg ctg cct gta ttt gaa agg aag ggc att gcg ctg Leu Gly Glu Val Leu Leu Pro Val Phe Glu Arg Lys Gly Ile Ala Leu 10 391 ggc aaa gtg gac atc tac ctg gac cag tcc aac aca ccc ctg tcc ctc Gly Lys Val Asp Ile Tyr Leu Asp Gln Ser Asn Thr Pro Leu Ser Leu 30 ace the gag gee tae agg the ggg gga cae tae ett egt gte aaa gee 439 Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr Leu Arg Val Lys Ala 487 cca gcc aag cct gga gat gag ggc aag gtg gag cag ggc atg aag gac Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu Gln Gly Met Lys Asp 60 535 tee aag tee etg agt ttg eeg att etg egg eea get ggg ace ggg eec Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro Ala Gly Thr Gly Pro

Pro Pro Pro Ser Pro Gln Pro Ala Ala Phe Pro Gln Ser Leu Pro

645

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				agt Ser												775
				ggc Gly 170												823
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				gcc Ala												967
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Thr Pro Leu Ser Leu Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr 35 40 45

Leu Arg Val Lys Ala Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu 50 55 60

Gln Gly Met Lys Asp Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro 65 70 75 80

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Arg Glu Ser Leu Asp Ile Leu Ala Pro Gly Arg Arg Arg Lys Asn Met 100 105 110

Ser Glu Phe Leu Gly Glu Ala Ser Ile Pro Gly Glu Glu Pro Pro Thr 115 120 125

Pro Ser Ser Cys Ser Leu Pro Ser Gly Ser Ser Gly Ser Thr Asn Thr 130 135 140

Gly Asp Ser Trp Lys Asn Arg Ala Ala Ser Arg Phe Ser Gly Phe Phe 145 150 155 160

Ser Ser Gly Pro Ser Thr Ser Ala Phe Gly Arg Glu Val Asp Lys Met 165 170 175

Glu Gln Leu Glu Gly Lys Leu His Thr Tyr Ser Leu Phe Gly Leu Pro 180 185 190

Arg Leu Pro Arg Gly Leu Arg Phe Asp His Asp Ser Trp Glu Glu Glu 195 200 205

Tyr Asp Glu Asp Glu Asp Glu Asp Asn Ala Cys Leu Arg Leu Glu Asp 210 215 220

Ser Trp Arg Glu Leu Ile Asp Gly His Glu Lys Leu Thr Arg Arg Gln 225 230 235 240

Cys His Gln Gln Glu Ala Val Trp Glu Leu Leu His Thr Glu Ala Ser 245 250 255

Tyr Ile Arg Lys Leu Arg Val Ile Ile Asn Leu Phe Leu Cys Cys Leu 260 265 270

Leu Asn Leu Gln Glu Ser Gly Leu Leu Cys Glu Val Glu Ala Glu Arg 275 280 285

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Gln Gln Leu Arg Ala Gln Glu Pro Pro Gly Ser Gln Gln Pro Leu Gln 615 Glu Glu Glu Glu Gly Glu Asp Ser Gly Thr Ser Ala Ala Ser Ser Pro Thr Ile Met Arg Lys Ser Ser Gly Ser Pro Asp Ser Gln His Cys 650 Ala Ser Asp Gly Ser Thr Glu Thr Leu Ala Met Val Val Val Glu Pro 665 Gly Asp Thr Leu Ser Ser Pro Glu Phe Asp Ser Gly Pro Phe Ser Ser 680 Gln Ser Asp Glu Thr Ser Leu Ser Thr Thr Ala Ser Ser Ala Thr Pro 695 Thr Ser Glu Leu Leu Pro Leu Gly Pro Val Asp Gly Arg Ser Cys Ser Met Asp Ser Ala Tyr Gly Thr Leu Ser Pro Thr Ser Leu Gln Asp Phe 730 Val Ala Pro Gly Pro Met Ala Glu Leu Val Pro Arg Ala Pro Glu Ser 745 Pro Arg Val Pro Ser Pro Pro Pro Ser Pro Arg Leu Arg Arg Arg Thr Pro Val Gln Leu Leu Ser Cys Pro Pro His Leu Leu Lys Ser Lys Ser 775 Glu Ala Ser Leu Leu Gln Leu Leu Ala Gly Ala Gly Thr His Gly Thr Pro Ser Ala Pro Ser Arg Ser Leu Ser Glu Leu Cys Leu Ala Val Pro 810 Ala Pro Gly Ile Arg Thr Gln Gly Ser Pro Gln Glu Ala Gly Pro Ser Trp Asp Cys Arg Gly Ala Pro Ser Pro Gly Ser Gly Pro Gly Leu Val Gly Cys Leu Ala Gly Glu Pro Ala Gly Ser His Arg Lys Arg Cys Gly 855 Asp Leu Pro Ser Gly Ala Ser Pro Arg Val Gln Pro Glu Pro Pro Pro Gly Val Ser Ala Gln His Arg Lys Leu Thr Leu Ala Gln Leu Tyr Arg 885 890

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cgaggcctgt gacagcaagt tccacagcac c atg cat tat gat ggg cat gtc Met His Tyr Asp Gly His Val 1 5	172
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acc egg tea tge eeg eeg ege acc age eee gea gtg gae ttg gag gag Thr Arg Ser Cys Pro Pro Arg Thr Ser Pro Ala Val Asp Leu Glu Glu 25 30 35	268
gag gag gag gag agc tct gtg gat ggc aaa ggg gac cgg aag agc aca Glu Glu Glu Glu Ser Ser Val Asp Gly Lys Gly Asp Arg Lys Ser Thr 40 45 50 55	316
ggc ctg aaa ctc tcc aag aag aaa gca agg agg aga cac acg gat gac Gly Leu Lys Leu Ser Lys Lys Lys Ala Arg Arg Arg His Thr Asp Asp 60 65 70	364
cca age aag gaa tge tte act etg aaa ttt gae etg aat gtg gae att Pro Ser Lys Glu Cys Phe Thr Leu Lys Phe Asp Leu Asn Val Asp Ile 75 80 85	412
gag aca gag atc gtc cca gcc atg aag aag aag tca ctg ggg gag gtg Glu Thr Glu Ile Val Pro Ala Met Lys Lys Lys Ser Leu Gly Glu Val 90 95 100	460
ctg ctg cct gta ttt gaa agg aag ggc att gcg ctg ggc aaa gtg gac Leu Leu Pro Val Phe Glu Arg Lys Gly Ile Ala Leu Gly Lys Val Asp 105 110 115	-508
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	Leu													ctg Leu		1276		

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caç Glr	ctg Leu 585	Leu	g ctg Leu	g gag ı Glu	ı Gly	ago Ser 590	Leu	g agg i Arg	g atg g Met	aaç Lys	gag Glu 595	Gly	aag Lys	gac Asp	agc Ser	1948

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ccc cgt ctc Pro Arg Leu						2764
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cct cag gaa Pro Gln Glu 920						2956
ggc agc ggt Gly Ser Gly						3004
tcc cac agg Ser His Arg	aag agg tgt Lys Arg Cys 955	gga gac cto Gly Asp Leo 960	ı Pro Ser	ggg gcc tct Gly Ala Ser 965	Pro Arg	3052
gtc cag cct Val Gln Pro 970						3100
acc ctg gcc Thr Leu Ala 985	cag ctc tac Gln Leu Tyr	cga atc ago Arg Ile Arc 990	g acc acc g Thr Thr	ctg ctg ctt Leu Leu Leu 995	aac tcc Asn Ser	3148
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Lys Gly Asp Arg Lys Ser Thr Gly Leu Lys Leu Ser Lys Lys Ala
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Arg Arg Arg His Thr Asp Asp Pro Ser Lys Glu Cys Phe Thr Leu Lys 65 70 75 80

Phe Asp Leu Asn Val Asp Ile Glu Thr Glu Ile Val Pro Ala Met Lys 85 90 95

Lys Lys Ser Leu Gly Glu Val Leu Leu Pro Val Phe Glu Arg Lys Gly
100 105 110

Ile Ala Leu Gly Lys Val Asp Ile Tyr Leu Asp Gln Ser Asn Thr Pro 115 120 125

Leu Ser Leu Thr Phe Glu Ala Tyr Arg Phe Gly Gly His Tyr Leu Arg 130 135 140

Val Lys Ala Pro Ala Lys Pro Gly Asp Glu Gly Lys Val Glu Gln Gly 145 150 155 160

Met Lys Asp Ser Lys Ser Leu Ser Leu Pro Ile Leu Arg Pro Ala Gly
165 170 175

Thr Gly Pro Pro Ala Leu Glu Arg Val Asp Ala Gln Ser Arg Arg Glu 180 185 190

Ser Leu Asp Ile Leu Ala Pro Gly Arg Arg Arg Lys Asn Met Ser Glu 195 200 205

Phe Leu Gly Glu Ala Ser Ile Pro Gly Gln Glu Pro Pro Thr Pro Ser 210 215 220

Ser Cys Ser Leu Pro Ser Gly Ser Ser Gly Ser Thr Asn Thr Gly Asp 225 230 235 240

Ser Trp Lys Asn Arg Ala Ala Ser Arg Phe Ser Gly Phe Phe Ser Ser

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Arg	Lys	Leu 355	Arg	Val	Ile	Ile	Asn 360	Leu	Phe	Leu	Cys	Cys 365	Leu	Leu	Asn
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Ser	Val	Leu	Arg 500	Lys	Thr	Glu	Glu	Pro 505		Ala	Lys	Glu	Ala 510	Val	Val
Ala	Met	Ile 515		Ser	Val	Glu	Arg 520		Ile	His	His	Val 525	Asn	Ala	Суѕ
Met	Arg 530		Arg	Gln	Glu	Arg 535	Gln	Arg	Leu	Ala	Ala 540	Val	Val	Ser	Arç

Ile Asp Ala Tyr Glu Val Val Glu Ser Ser Ser Asp Glu Val Asp Lys

545					550					555					560
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Met	Lys	Glu 595	Gly	Lys	Asp	Ser	Lys 600	Met	Asp	Val	Tyr	Cys 605	Phe	Leu	Phe
Thr	Asp 610	Leu	Leu	Leu	Val	Thr 615	Lys	Ala	Val	Lys	Lys 620	Ala	Glu	Arg	Thr
Arg 625	Val	Ile	Arg	Pro	Pro 630	Leu	Ļeu	Val	Asp	Lys 635	Ile	Val	Cys	Arg	Glu 640
Leu	Arg	Asp	Pro	Gly 645	Ser	Phe	Leu	Leu	Ile 650	Tyr	Leu	Asn	Glu	Phe 655	His
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Ile	Met	Arg	Lys 740	Ser	Ser	Gly	Ser	Pro 745	Asp	Ser	Gln	His	Cys 750	Ala	Ser
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Thr	Leu 770	Ser	Ser	Pro	Glu	Phe 775		Ser	Gly	Pro	Phe 780	Ser	Ser	Gln	Ser
Asp 785		Thr	Ser	Leu	Ser 790		Thr	Ala	Ser	Ser 795	Ala	Thr	Pro	Thr	Ser 800
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Ser Val Asn Ala Glu Asp Glu Glu Gly Asp Thr Ala Leu His Val Ala
740 745 750

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												tgg Trp				2129
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	gta Val	_	_		_	_	_				-		tga	tctg	gcg	2658
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gga	aggga	atg :	tgct	gttt	cc ct	cta	egtge	c cca	agtco	ctag	cct	cgct	cta	ggac	ccaggg	2778.
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Asn Ile Tyr Thr Phe Asn His Thr Val Thr Arg Asn Arg Thr Glu Gly
Val Arg Val Ser Val Asn Val Leu Asn Lys Gln Lys Gly Ala Pro Leu
Leu Phe Val Val Arg Gln Lys Glu Ala Val Val Ser Phe Gln Val Pro
Leu Ile Leu Arg Gly Met Phe Gln Arg Lys Tyr Leu Tyr Gln Lys Val
Glu Arg Thr Leu Cys Gln Pro Pro Thr Lys Asn Glu Ser Glu Ile Gln
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Gln Leu Arg Val Ser Arg Met Asp Asp Phe Val Leu Arg Thr Gly Glu
145
                    150
                                        155
Gln Phe Ser Phe Asn Thr Thr Ala Ala Gln Pro Gln Tyr Phe Lys Tyr
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Glu Phe Pro Glu Gly Val Asp Ser Val Ile Val Lys Val Thr Ser Asn
Lys Ala Phe Pro Cys Ser Val Ile Ser Ile Gln Asp Val Leu Cys Pro
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Val Tyr Asp Leu Asp Asn Asn Val Ala Phe Ile Gly Met Tyr Gln Thr
Met Thr Lys Lys Ala Ala Ile Thr Val Gln Arg Lys Asp Phe Pro Ser
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Asn Ser Phe Tyr Val Val Val Val Lys Thr Glu Asp Gln Ala Cys

Gly Gly Ser Leu Pro Phe Tyr Pro Phe Ala Glu Asp Glu Pro Val Asp 260 265 270

230

Gln Gly His Arg Gln Lys Thr Leu Ser Val Leu Val Ser Gln Ala Val Thr Ser Glu Ala Tyr Val Ser Gly Met Leu Phe Cys Leu Gly Ile Phe Leu Ser Phe Tyr Leu Leu Thr Val Leu Leu Ala Cys Trp Glu Asn Trp 310 315 Arg Gln Lys Lys Thr Leu Leu Val Ala Ile Asp Arg Ala Cys Pro 330 Glu Ser Gly His Pro Arg Val Leu Ala Asp Ser Phe Pro Gly Ser Ser 345 Pro Tyr Glu Gly Tyr Asn Tyr Gly Ser Phe Glu Asn Val Ser Gly Ser 360 Thr Asp Gly Leu Val Asp Ser Ala Gly Thr Gly Asp Leu Ser Tyr Gly Tyr Gln Gly His Asp Gln Phe Lys Arg Arg Leu Pro Ser Gly Gln Met Arg Gln Leu Cys Ile Ala Met Gly Arg Ser Phe Glu Pro Val Gly Thr 410 Arg Pro Arg Val Asp Ser Met Ser Ser Val Glu Asp Asp Tyr Asp Thr Leu Thr Asp Ile Asp Ser Asp Lys Asn Val Ile Arg Thr Lys Gln Tyr Leu Tyr Val Ala Asp Leu Ala Arg Lys Asp Lys Arg Val Leu Arg Lys Lys Tyr Gln Ile Tyr Phe Trp Asn Ile Ala Thr Ile Ala Val Phe Tyr Ala Leu Pro Val Val Gln Leu Val Ile Thr Tyr Gln Thr Val Val Asn Val Thr Gly Asn Gln Asp Ile Cys Tyr Tyr Asn Phe Leu Cys Ala His Pro Leu Gly Asn Leu Ser Ala Phe Asn Asn Ile Leu Ser Asn Leu Gly Tyr Tle Leu Leu Gly Leu Leu Phe Leu Leu Ile Ile Leu Gln Arg 535 Glu Ile Asn His Asn Arg Ala Leu Leu Arg Asn Asp Leu Cys Ala Leu

545

570

Glu Cys Gly Ile Pro Lys His Phe Gly Leu Phe Tyr Ala Met Gly Thr

Ala Leu Met Met Glu Gly Leu Leu Ser Ala Cys Tyr His Val Cys Pro Asn Tyr Thr Asn Phe Gln Phe Asp Thr Ser Phe Met Tyr Met Ile Ala 600 Gly Leu Cys Met Leu Lys Leu Tyr Gln Lys Arg His Pro Asp Ile Asn 615 Ala Ser Ala Tyr Ser Ala Tyr Ala Cys Leu Ala Ile Val Ile Phe Phe 630 635 Ser Val Leu Gly Val Val Phe Gly Lys Gly Asn Thr Ala Phe Trp Ile 650 Val Phe Ser Ile Ile His Ile Ile Ala Thr Leu Leu Ser Thr Gln Leu Tyr Tyr Met Gly Arg Trp Lys Leu Asp Ser Gly Ile Phe Arg Arg Ile Leu His Val Leu Tyr Thr Asp Cys Ile Arg Gln Cys Ser Gly Pro Leu Tyr Val Asp Arg Met Val Leu Leu Val Met Gly Asn Val Ile Asn Trp Ser Leu Ala Ala Tyr Gly Leu Ile Met Arg Pro Asn Asp Phe Ala Ser Tyr Leu Leu Ala Ile Gly Ile Cys Asn Leu Leu Tyr Phe Ala Phe Tyr Ile Ile Met Lys Leu Arg Ser Gly Glu Arg Ile Lys Leu Ile Pro Leu Cys Ile Val Cys Thr Ser Val Val Trp Gly Phe Ala Leu Phe Phe Phe Gln Gly Leu Ser Thr Trp Gln Lys Thr Pro Ala Glu Ser Arg Glu His Asn Arg Asp Cys Ile Leu Leu Asp Phe Phe Asp Asp 810 His Asp Ile Trp His Phe Leu Ser Ser Ile Ala Met Phe Gly Ser Phe Leu Val Ser Gly Pro Pro Gly Arg Ala Gly Trp Val Arg Glu Gly Ser 840 Ser Cys Leu Leu Pro Cys Gly

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Glu Cys Thr Glu Ala Lys Lys His Cys Trp Tyr Phe Glu Gly Leu Tyr
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Pro Thr Tyr Tyr Ile Cys Arg Ser Tyr Glu Asp Cys Cys Gly Ser Arg
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Cys Cys Val Arg Ala Leu Ser Ile Gln Arg Leu Trp Tyr Phe Trp Phe
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Arg Arg Met Tyr Pro Pro Pro Leu Ile Glu Glu Pro Ala Phe Asn
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                                                                   394
Val Ser Tyr Thr Arg Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln Gln
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ccq qqq ccg ccc tat tac acc gac cca gga gga ccg ggg atg aac cct
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Pro Gly Pro Pro Tyr Tyr Thr Asp Pro Gly Gly Pro Gly Met Asn Pro
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gtc ggg aat tcc atg gca atg gct ttc cag gtc cca ccc aac tca ccc
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Val Gly Asn Ser Met Ala Met Ala Phe Gln Val Pro Pro Asn Ser Pro
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                                        140
cag ggg agt gtg gcc tgc ccg ccc cct cca gcc tac tgc aac acg cct
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Gln Gly Ser Val Ala Cys Pro Pro Pro Pro Ala Tyr Cys Asn Thr Pro
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ccg ccc ccg tac gaa cag gta gtg aag gcc aag tagtggggtg cccacgtgca 591
Pro Pro Pro Tyr Glu Gln Val Val Lys Ala Lys
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Arg Cys Cys Val Arg Ala Leu Ser Ile Gln Arg Leu Trp Tyr Phe Trp 50 60

Phe Leu Leu Met Met Gly Val Leu Phe Cys Cys Gly Ala Gly Phe Phe 65 70 75 80

Ile Arg Arg Met Tyr Pro Pro Pro Leu Ile Glu Glu Pro Ala Phe
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Asn Val Ser Tyr Thr Arg Gln Pro Pro Asn Pro Gly Pro Gly Ala Gln
100 105 110

Gln Pro Gly Pro Pro Tyr Tyr Thr Asp Pro Gly Gly Pro Gly Met Asn 115 120 125

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														aca Thr		1707
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Ser Gly Asn Arg Asp Thr Leu Trp His Leu Phe Asn Thr Leu Gln Arg 50 55 60

Arg Pro Gly Trp Val Glu Tyr Phe Ile Ala Ala Leu Arg Gly Cys Glu

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Pro	Tyr 130	Asn	Ser	Cys	Arg	Glu 135	Lys	Glu	Pro	Ser	Tyr 140	Pro	Met	Pro	Val
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Leu	Gln	Thr	Leu	Ser 165	Pro	Arg	Ala	Ile	Pro 170	Arg	Asn	Pro	Asp	Gly 175	Gly
Pro	Leu	Glu	Ser 180	Ser	Ser	Asp	Leu	Ala 185	Ala	Leu	Ser	Pro	Leu 190	Thr	Ser
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Ser	Thr	Arg 355	Ala	Gly	Met	Val	Pro 360	Ser	Lys	Val	Pro	Thr 365	Ser	Met	Val
Leu	Thr	Lys	Val	Ser	Ala	Ser	Thr	Val	Pro	Thr	Asp	Glv	Ser	Ser	Ara

370 375 380 Asn Glu Glu Thr Pro Ala Ala Pro Thr Pro Ala Gly Ala Thr Gly Gly 385 390 395 Ser Ser Ala Trp Leu Asp Ser Ser Ser Glu Asn Arg Gly Leu Gly Ser 405 410 415 Glu Leu Ser Lys Pro Gly Val Leu Ala Ser Gln Val Asp Ser Pro Phe 425 420 Ser Gly Cys Phe Glu Asp Leu Ala Ile Ser Ala Ser Thr Ser Leu Gly 440 Met Gly Pro Cys His Gly Pro Glu Glu Asn Glu Tyr Lys Ser Glu Gly Thr Phe Gly Ile His Val Ala Glu Asn Pro Ser Ile Gln Leu Leu Glu 470 475 Gly Asn Pro Gly Pro Pro Ala Asp Pro Asp Gly Gly Pro Arg Pro Gln 490 Ala Asp Arg Lys Phe Gln Glu Arg Glu Val Pro Cys His Arg Pro Ser Pro Gly Ala Leu Trp Leu Gln Val Ala Val Thr Gly Val Leu Val Val Thr Leu Leu Val Val Leu Tyr Arg Arg Arg Leu His 530 535 <210> 171 <211> 3579 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (242)..(3094)

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														gac Asp 95		529
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Lys Gly Pro Leu Pro Met Glu Ala Ile Glu Lys Met Ala Ser Leu Cys
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Pro Glu Ala Pro His Pro Gly Leu Glu Thr Thr Leu Gln Glu Arg Leu 130 135 140 Ala Leu Tyr Gln Thr Ala Ile Glu Ser Ala Arg Gln Ala Gly Asp Ser Ala Lys Met Arg Arg Tyr Asp Arg Gly Leu Lys Thr Leu Glu Asn Leu 170 Leu Ala Ser Ile Arg Lys Gly Asn Ala Ile Asp Glu Ala Asp Ile Pro 185 Pro Pro Val Ala Ile Gly Lys Gly Pro Ala Ser Thr Pro Thr Tyr Ser 200 Pro Ala Pro Thr Gln Pro Ala Pro Arg Ile Ala Ser Ala Pro Glu Pro 215 Arg Val Thr Leu Glu Gly Pro Ser Ala Thr Ala Pro Ala Ser Ser Pro 235 Gly Leu Ala Lys Pro Gln Met Pro Pro Gly Pro Cys Ser Pro Gly Pro Leu Ala Gln Leu Gln Ser Arg Gln Arg Asp Tyr Lys Leu Ala Ala Leu His Ala Lys Gln Gln Gly Asp Thr Thr Ala Ala Ala Arg His Phe Arg Val Ala Lys Ser Phe Asp Ala Val Leu Glu Ala Leu Ser Arg Gly Glu Pro Val Asp Leu Ser Cys Leu Pro Pro Pro Pro Asp Gln Leu Pro Pro 310 Asp Pro Pro Ser Pro Pro Ser Gln Pro Pro Thr Pro Ala Thr Ala Pro 325 330 Ser Thr Thr Glu Val Pro Pro Pro Pro Arg Thr Leu Leu Glu Ala Leu 345 Glu Gln Arg Met Glu Arg Tyr Gln Val Ala Ala Ala Gln Ala Lys Ser Lys Gly Asp Gln Arg Lys Ala Arg Met His Glu Arg Ile Val Lys Gln 375 Tyr Gln Asp Ala Ile Arg Ala His Lys Ala Gly Arg Ala Val Asp Val Ala Glu Leu Pro Val Pro Pro Gly Phe Pro Pro Ile Gln Gly Leu Glu 410 Ala Thr Lys Pro Thr Gln Gln Ser Leu Val Gly Val Leu Glu Thr Ala Met Lys Leu Ala Asn Gln Asp Glu Gly Pro Glu Asp Glu Glu Asp Glu 440

Val Pro Lys Lys Gln Asn Ser Pro Val Ala Pro Thr Ala Gln Pro Lys Ala Pro Pro Ser Arg Thr Pro Gln Ser Gly Ser Ala Pro Thr Ala Lys 470 Ala Pro Pro Lys Ala Thr Ser Thr Arg Ala Gln Gln Gln Leu Ala Phe 485 490 Leu Glu Gly Arg Lys Lys Gln Leu Leu Gln Ala Ala Leu Arg Ala Lys 505 Gln Lys Asn Asp Val Glu Gly Ala Lys Met His Leu Arg Gln Ala Lys 520 Gly Leu Glu Pro Met Leu Glu Ala Ser Arg Asn Gly Leu Pro Val Asp Ile Thr Lys Val Pro Pro Ala Pro Val Asn Lys Asp Asp Phe Ala Leu Val Gln Arg Pro Gly Pro Gly Leu Ser Gln Glu Ala Ala Arg Arg Tyr 570 Gly Glu Leu Thr Lys Leu Ile Arg Gln Gln His Glu Met Cys Leu Asn His Ser Asn Gln Phe Thr Gln Leu Gly Asn Ile Thr Glu Thr Thr Lys 595 Phe Glu Lys Leu Ala Glu Asp Cys Lys Arg Ser Met Asp Ile Leu Lys Gln Ala Phe Val Arg Gly Leu Pro Thr Pro Thr Ala Arg Phe Glu Gln 630 635 Arg Thr Phe Ser Val Ile Lys Ile Phe Pro Asp Leu Ser Ser Asn Asp 650 Met Leu Leu Phe Ile Val Lys Gly Ile Asn Leu Pro Thr. Pro Pro Gly Leu Ser Pro Gly Asp Leu Asp Val Phe Val Arg Phe Asp Phe Pro Tyr 680 Pro Asn Val Glu Glu Ala Gln Lys Asp Lys Thr Ser Val Ile Lys Asn Thr Asp Ser Pro Glu Phe Lys Glu Gln Phe Lys Leu Cys Ile Asn Arg 710 715 Ser His Arg Gly Phe Arg Arg Ala Ile Gln Thr Lys Gly Ile Lys Phe Glu Val Val His Lys Gly Gly Leu Phe Lys Thr Asp Arg Val Leu Gly 740 745

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Glu Val Met Val Arg Ile Arg Glu Pro Leu Thr Ala Gln Gln Leu Glu
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Thr Thr Glu Arg Trp Leu Val Ile Asp Pro Val Pro Ala Ala Val
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Pro Thr Gln Val Ala Gly Pro Lys Gly Lys Ala Pro Pro Val Pro Ala
Pro Ala Arg Glu Ser Gly Asn Arg Ser Ala Arg Pro Leu His Ser Leu
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Ser Val Leu Ala Phe Asp Gln Glu Arg Leu Glu Arg Lys Ile Leu Ala
Leu Arg Gln Ala Arg Arg Pro Val Pro Pro Glu Val Ala Gln Gln Tyr
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Gln Asp Ile Met Gln Arg Ser Gln Trp Gln Arg Ala Gln Leu Glu Gln
Gly Gly Val Gly Ile Arg Arg Glu Tyr Thr Ala Gln Leu Glu Arg Gln
            900
Leu Gln Phe Tyr Thr Glu Ala Ala Arg Arg Leu Gly Asn Asp Gly Ser
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Val Phe Ile Leu Gly Leu Ile Thr Asn Ser Ala Ser Leu Phe Val Phe 50 60

Cys Phe Arg Met Lys Met Arg Ser Glu Thr Ala Thr Phe Ile Thr Asn 65 70 75 80

Leu Ala Leu Ser Asp Leu Leu Phe Val Cys Thr Leu Pro Phe Lys Ile 85 90 95

Phe Tyr Asn Phe Asn Arg His Trp Pro Phe Gly Asp Thr Leu Cys Lys 105 Ile Ser Gly Thr Ala Phe Leu Thr Asn Ile Tyr Gly Ser Met Leu Phe 120 Leu Thr Cys Ile Ser Val Asp Arg Phe Leu Ala Ile Val Tyr Pro Phe Arg Ser Arg Thr Ile Arg Thr Arg Arg Asn Ser Ala Ile Val Cys Ala Gly Val Trp Ile Leu Val Leu Ser Gly Gly Ile Ser Ala Ser Leu Phe 165 Ser Thr Thr Asn Val Asn Asn Ala Thr Thr Thr Cys Phe Glu Gly Phe 180 185 Ser Lys Arg Val Trp Lys Thr Tyr Leu Ser Lys Ile Thr Ile Phe Ile Glu Val Val Gly Phe Ile Ile Pro Leu Ile Leu Asn Val Ser Cys Ser 215 Ser Val Val Leu Arg Thr Leu Arg Lys Pro Ala Thr Leu Ser Gln Ile Gly Thr Asn Lys Lys Lys Val Leu Lys Met Ile Thr Val His Met Ala 250 Val Phe Val Cys Phe Val Pro Tyr Asn Ser Val Leu Phe Leu Tyr Ala Leu Val Arg Ser Gln Ala Ile Thr Asn Cys Leu Leu Glu Arg Phe Ala Lys Ile Met Tyr Pro Ile Thr Leu Cys Leu Ala Thr Leu Asn Cys Cys Phe Asp Pro Phe Ile Tyr Tyr Phe Thr Leu Glu Ser Phe Gln Lys 305 310 Ser Phe Tyr Ile Asn Thr His Ile Arg Met Glu Ser Leu Phe Lys Thr 330 Glu Thr Pro Leu Thr Pro Lys Pro Ser Leu Pro Ala Ile Gln Glu Glu Val Ser Asp Gln Thr Thr Asn Asn Gly Gly Glu Leu Met Leu Glu Ser 365

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Val Phe Ile Leu Gly Leu Ile Thr Asn Ser Val Ser Leu Phe Val Phe 50 55 60

Cys Phe Arg Met Lys Met Arg Ser Glu Thr Ala Ile Phe Ile Thr Asn 65 70 75 80

Leu Ala Val Ser Asp Leu Leu Phe Val Cys Thr Leu Pro Phe Lys Ile $85 \ 90 \ 95$

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Ala	Leu	Val 275	Arg	Ser	Gln	Ala	Ile 280	Thr	Asn	Cys	Phe	Leu 285	Glu	Arg	Phe
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Val Ala Gly Glu Gln Ala Pro Gly Thr Ala Pro Cys Ser Arg Gly Ser
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Ser Trp Ser Ala Asp Leu Asp Lys Cys Met Asp Cys Ala Ser Cys Arg
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Ala Arg Pro His Ser Asp Phe Cys Leu Gly Cys Ala Ala Ala Pro Pro
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Gly Cys Pro Ala Val Ala Leu Ile Gln
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Cys Met Asp Cys Ala Ser Cys Arg Ala Arg Pro His Ser Asp Phe Cys 50 60

Leu Gly Cys Ala Ala Ala Pro Pro Ala Pro Phe Arg Leu Leu Trp Pro 65 70 75 80

Ile Leu Gly Gly Ala Leu Ser Leu Thr Phe Val Leu Gly Leu Leu Ser 85 90 95

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